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Run
                                                                                                                                                      Minimum DB
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                                                                                                                                                                                                                                                                Sequence:
                                                                                                                                                                                                                                                                           Title:
Perfect score:
                                                                                                                                                                                                                                                                                                                                                      OM protein
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                                                                                                                     Post-processing: Minimum Match
Maximum Match
                                                                                                                                                                                    Total number of hits satisfying chosen parameters:
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                                                                                                                                                     seq length: 0 seq length: 40
                                                                                                                                                                                                                                                                                                                                                   protein search, using sw model
BLOSUM62
Gapop 10.0 , Gapext 0.5
                                                                                                                                                                                                                                                                            US-09-509-449C-5
123
                                                                                                                                                                                                                                                                 1 PRGSRPSWGPTDPRHRSRNVG
                                                                                                                                                                                                                                                                                                                                June
                                                                                                                                                                                                          1586107 seqs, 282547505 residues
                                                                                                        Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                                                                                                                                                                                                                                                                                                                                                     Copyright
                                                                                                                                                                                                                                                                                                         GenCore version 5.1.6 (c) 1993 - 2004 Compugen Ltd
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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r55582	518	615	j03266 Hepatiti	03656 Hepatit	j03571 Hepatiti	30986 Sho	Aau84605 HCV HepCl	4515	7126 Control	0830 HCV	Aar31922 HCV pepti	4604 HCV	56608 HCV	NANB	mok	Aar67636 Epitope o	4 NANB	Aar38293 NANB hepa	347 P99 HCV	33863 Po.	1 HCV-	34055 HCV	O He		Description		

A method has been developed for virus-containing sample is to use a treatment solution comprising: (i) an anionic surfactant; (ii) an amphoteric surfactant; (iii) a non-ionic surfactant; and (iv) a protein denaturing agent. Also described in the present invention are: (1) a similar treatment method by using a solution containing: (i) a chaotropic ion; (ii) an acidifying agent; and (iii) a non-ionic surfactant; (2) an assaying method for viruses by quantitative detection of the presence of viral antigen using a probe for reaction with the specifically recognised viral antigen; (3) a hybridoma cell line chosen from HC11-1 (FERM BP-6005), HC11-14 (FERM BP-6006), HC11-0 (FERM BP-6004), HC11-3 (FERM BP-6002) and HC11-7 (FERM BP-6003); (4) a monoclonal antibody produced by

Detection and assay for, e.g. hepatitis C virus - comprises treatment the sample with surfactants, protein denaturing agent prior to

of.

Example 3; Page 88; 98pp; Japanese.

immunoassay.

<b>4</b> UI	44	<b>4</b> ω	42	41	40	39	38	37	36	ယ ပၢ	34	ω ω	32	31	30	29	28	27	26
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AAJ00501	AAJ01576	AAJ00986	AAJ02919	AAJ03613	AAJ03221	AAJ02595	N	AAY80191	AAW99649	AAR30685	AAR84400	AAY44550	AAJ02596	AAJ02920	AAJ03466	AAW10269	ABR55583	AAR56607	AAJ03982
Aaj00501	Aaj01576	Aaj00986	Aaj02919	Aaj03613	Aaj03221	Aaj02595	Aaj 03528	Aay80191	Aaw99649	Aar30685	Aar84400	Aay44550	Aaj02596	Aaj02920	Aaj03466	Aaw10269	Abr55583	Aar56607	Aaj03982
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### ALIGNMENTS

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04-AUG-1997;
04-AUG-1997;
31-JUL-1998;
                                                                                                                                                                                             Hepatitis C virus; HCV; hepatitis B virus; diagnosis; detection viral infection.
                                                                                               WPI; 1999-153953/13.
                                                                                                         Aoyagi K,
                                                                                                                                                   04-AUG-1998;
                                                                                                                                                                                   Hepatitis C virus.
                                                                                                                                                                                                             Hepatitis C virus peptide SEQ ID NO:5.
                                                                                                                                                                                                                        21-MAY-1999
                                                                                                                                                                                                                                  AAW99648;
                                                                                                                                                                                                                                            AAW99648 standard; peptide;
                                                                                                                                                              11-FEB-1999.
                                                                                                                                                                         WO9906836-A1.
                                                                                                                    (TOFU ) TONEN CORP
                                                                                                        Ohue C,
                                                                                                                                                                                                                       (first entry)
                                                                                                                             97JP-00209515.
97JP-00209522.
98JP-00218136.
                                                                                                                                                   98WO-JP003476.
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                                                                                                         Kimura
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                                                                                                         Yagi
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RESULT 2
AAY80190
ID AAY8
AXX AAY8
AC AAY8
AC AAY8
AC AAY8
AXX Hepa
XXX AO91
IDR W020
PPT W1XI
PPT Met]
PPT Met
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local S
Matches 21
A method has been developed of assaying for hepatitis C virus (HCV). The method comprises binding a viral core antigen or anti-HCV antibody to a suitable probe in the presence of a non-ionic surfactant and/or a cationic surfactant with secondary, tertiary or quaternary amine groups and carbon chains with 10 or more carbon atoms. The assay is useful for the detection of HCV infection in biological samples, especially in donated blood. The method is highly sensitive for detecting HCV and reduces the window period between negative and positive screening tests for HCV on blood donated by a recently infected person. The present sequence represents an HCV peptide which is used in an example from the present invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     screening
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Hepatitis C virus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Hepatitis C
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 3; Page 45; 55pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             the presence of a surfactant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Method of assaying for hepatitis C virus infection comprises binding viral core antigen or anti-hepatitis C virus antibodies with a probe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2000-195352/17.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               30-JUL-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     30-JUL-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Aoyagi K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (ADLI-) ADVANCED LIFE SCI INST INC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Ohue C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              virus peptide sequence SEQ ID NO:5.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    HCV; antigen; antibody; infection; blood donation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                peptide;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      100.0%;
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Pred. No. 3.2e-10;
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RESULT 4
ABJ37517
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AC ABJ3
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Best Local (
                                                                                                                                                           Best
                                                                                                                                                                                                                       The synthetic peptide corresponds to residues 98-121 of the C domain of HCV. The peptide may be used for the sensitive and specific detection of HCV infection Antibodies raised against the peptide can be used for passive immuno-prophylaxis and the HCV antigens can be used in vaccines to prevent HCV infection. See also AAR34041-65. (Updated on 25-MAR-2003)
                                                                                                                                                                                                            passive immuno-prophylaxis and the to prevent HCV infection. See also to correct PN field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               25-MAR-2003
22-JUL-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence
           10-MAY-2003
                                 ABJ37517;
                                                    ABJ37517
                                                                                                                                                                                        Sequence 24
                                                                                                                                                                                                                                                                                    Example 7; Page 130; 190pp; English.
                                                                                                                                                                                                                                                                                                        Detection of hepatitis C virus antigens - using HCV reactive antibody bound to solid support and competitive HCV antigen-reporter complex.
                                                                                                                                                                                                                                                                                                                                         WPI; 1993-117737/14.
                                                                                                                                                                                                                                                                                                                                                             Dreesman GR,
                                                                                                                                                                                                                                                                                                                                                                                                     16-SEP-1991;
12-FEB-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                    16-SEP-1992;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Hepatitis C virus; non structural region; antigen; diagnosis; vaccine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           HCV C domain
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                                                    standard;
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                                                                                                                                                 Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           peptide DP10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (revised)
(first entry)
           (first
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                                                                                                                                                                                                                                                                                                                                                                                                      91US-00762135.
92US-00835717.
                                                                                                                                                                                                                                                                                                                                                                                                                                    92WO-US007865
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  peptide; 24
                                                    peptide;
           entry)
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                                                                                                                                                          91.1%;
                                                                                                                                                                                                                                                                                                                                                             Pauletti D;
                                                     25
                                                                                                                                                Score 112; DB Pred. No. 1.2e 1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0,
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Pred. No. 3.2e-10;
'; Mismatches 0;
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                                                                                                                                                          DB 2;
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RESULT 5
AAR33863
ID AAR3
XX AAR3
AC AAR3
AC AAR3
XX 19-1
CX PO1:
XX P40
XX P40
XX P40
XX Syn
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PP 16:
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                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local S
Matches 19
                                                                                                                                                                                                                                                                                                                                                                                                             The invention relates to a novel method for simultaneously detecting a hepatitis C virus (HCV) antigen and an HCV antibody in a test sample. The novel method comprises contacting the test sample with the antigen and antibody, both coated on solid phases, independently, for a time and under conditions sufficient for the formation of an antibody/antigen complex and a second antibody/antigen complex, and detecting the two complexes, which indicate the presence of the antigen and antibody in the test sample. The method is useful for detecting a hepatitis C virus (HCV) antigen and HCV antibody in a test sample. This sequence represents an HCV-core derived peptide relating to the HCV antigen/antibody detection method of the invention
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Shah DO,
Desai S,
                                                                                                   Hepatitis p408.
                                                                                                                                                             25-MAR-2003
19-JUL-1993
                                                                                                                                                                                                                                                                                                                                                                                        Sequence 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Detecting hepatitis C virus antigen and antibody in sample, by con sample with HCV antigen and antibody, for forming HCV antigen-test antibody complex and HCV antibody-test antigen complex, detecting
 16-SEP-1992;
                         01-APR-1993.
                                                                          Synthetic
                                                                                                                                     Polypeptide
                                                                                                                                                                                                                         AAR33863 standard; peptide; 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 1; Page 17; 92pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          complexes.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2003-201507/19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                24-JUN-2002;
                                                  WO9306247-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             26-JUN-2001;
17-JUN-2002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Hepatitis C virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Hepatitis C virus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (ABBO ) ABBOTT LAB.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            HCV-core
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                                                                                                                                                                                                                                                                                                                                                    Similarity
                                                                                                                                                                                                                                                                                      PRGSRPSWGPTDPRRRSRNLG
                                                                                                                                                                                                                                                                                                            PRGSRPSWGPTDPRHRSRNVG 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          derived peptide
                                                                                                             a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Dawson GA,
Stewart JL;
                                                                                                                                    66d
                                                                                                                                                                                                                                                                                                                                                                                        AA;
                                                                                                           virus; NANBH; assay; antibody; p380-JH1; p380-J; p380LG;
                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                             (revised)
(first en
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2001US-00891983
2002US-00173480
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2002WO-US019958
 92WO-US007813
                                                                                                                                    comprising HCV viral antigen.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  HCV; antigen; antibody.
                                                                                                                                                             entry)
                                                                                                                                                                                                                                                                                                                                                 91.1%;
90.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Muerhoff
                                                                                                                                                                                                                                                                                                                                                   Score 112; DB 6;
Pred. No. 1.3e-08;
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                                                                                                                                                                                                                                                                                       25
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                                                                                                                                                                                                                                                                                                                                        Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Jiang
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ۲
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gutierrez
                                                                                                                                                                                                                                                                                                                                                            Length 25
                                                                                                                                                                                                                                                                                                                                       Indels
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                                                                                                                                                                                                                                                                                                                                     Gaps
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RESULT 6
AAR13347
ID AAR13347
ID 25-y
DT 25-y
DT 23-C
XX C100
XX C100
XX Synt
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XX AU90
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Best Local S
Matches 19
                                            The polypeptide may be prepared by solid phase synthesis fragment coupling (pref.) or using recombinant technology. The assay has increased sensitivity and is more specific than assays using the polypeptide C100-3 (EP-318216). See also AAQ13146-48 and AAR13343-65. (Updated on 25-MAR-2003 to correct PA field.)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The synthetic peptide p99 represents amino acid residues 99-126 of the hepatitis C viral antigen. The peptide may be used in an assay to detec antibodies to HCV and thus to diagnose chronic HCV infection. See also AAR33862-87. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                                                                                Immunological assays for hepatitis C virus antibody - by using polypeptide(s) contg. epitope(s) of hepatitis C virus antigens
                                                                                                                                                                                                                                                                                                                             WPI; 1991-238393/33.
                                                                                                                                                                                                                                                                                                                                                                               (ABBO )
                                                                                                                                                                                                                                                                                                                                                                                                                                                    22-DEC-1989;
07-NOV-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 C100-3; hepatitis C virus; immunoassay; epitope.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                25-MAR-2003
23-OCT-1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAR13347 standard;
                                                                                                                                                                                                   Claim 10; Page 48;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                21-DEC-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                27-JUN-1991.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              P99 HCV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAR13347;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure; Page 12; 63pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Assay for detecting
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Lesniewski RR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      16-SEP-1991;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  say for detecting presence of antibody to hepatitis C viral antigen - contacting sample with polypeptide contg. at least one epitope of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 PRGSRPSWGPTDPRHRSRNVG
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LESNIEWSKI
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (revised)
(first ent
                                                                                                                                                                                                                                                                                                                                                                                                                                                    89US-00456162
90US-00610180
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                90AU-00068390
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                                                                                                                                                                                                62pp; English.
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Pred. No. 1
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.4e-08;
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Seguence 29

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Best Local S
Matches 19
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Best Local S
Matches 19
                                                                                                                                                                                                                                                                                                                                                      method
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                            NANB hepatitis virus HC-OM gene
                                                                                                                                                                                                                                                                                                          Sequence 20
                                                                                                                                                                                                                                                                                                                                      The sequence is that of NANB hepatitis virus HC-OM gene polypeptide CP-5-2. It may may be used in a system for detecting NANB hepatitis. This method is highly specific and sensitive, and can detect NANB hepatitis virus which could not be detected by conventional methods
                                                                                                                                                                                                                                                                                                                                                                                                                                        Antigen related to non-A and non-B hepatitis virus - comprises non-translation region comprising 340 - 341 mols. of nucleotides, non-translation region comprising 1885 - 2551 mols. of nucleotides included non-translation region 1,149 and, etc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         12-JUN-1990;
08-NOV-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              JP05091884-A
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                                                         21-OCT-1993
                                                                                                                                                                                                                                                                                                                                                                                                              Claim 29; Page 63; 73pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     10-APR-1991;
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                                                                                       AAR38294;
                                                                                                                   AAR38294 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 N-PSDB; AAQ43903.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 1993-199637/25.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              hepatitis virus HC-OM
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ic; HCV; NANBH
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19; Conserv
non-B;
                                                                                                                                                                                                                      N
                                                                                                                                                                                                                                                                  Similarity
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                                                                                                                                                                                          RGSRPSWGPNDPRHRSRNVG
                                                                                                                                                                                                                                                                                                            ΑA;
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                                                           (first
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virus; polymerase
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90JP-00304405.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      91JP-00196175.
                                                                                                                 protein;
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                                                         entry)
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                                                                                                                                                                                                                                                               90.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  polymerase chain reaction; detection; sensitive;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                gene
                                                                                                                                                                                                                                               Score 111; DB 2; Le
Pred. No. 1.4e-08;
"" wismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 112; DB 2;
Pred. No. 1.5e-08;
1; Mismatches 1
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                            polypeptide CP-5-3.
 chain reaction;
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   detection;
                                                                                                                                                                                                                                                                              Length
                                                                                                                                                                                                                                                    Indels
                                                                                                                                                                                                                                                                                 20
 sensitive;
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   RESULT 9
AAR67636
ID AAR67636
AC AAR6

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Best Local S
Matches 18
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25-MAR-2003
18-AUG-1995
                                                                                                                                                                                            Non-A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Non-A.
non-B hepatitis
                                                           28-MAY-1993;
02-MAR-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 20 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            12-JUN-1990;
08-NOV-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                JP05091884-A
    Aoyama M,
                                                                                                        27-MAY-1994;
                                                                                                                                    14-DEC-1994
                                                                                                                                                                EP628572-A2
                                                                                                                                                                                                                                                                                               Epitope
                                                                                                                                                                                                                                                                                                                                                                                                                   AAR67636 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 30; Page
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                               (EISA ) EISAI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (NAKA/) NAKAMURA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1993-199637/25.
DB; AAQ43904.
                                                                                                                                                                                            hepatitis
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    Obara T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                             (revised)
(revised)
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      Tohmatsu
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Antigen related to non-A and non-B hepatitis virus - comprises non-translation region comprising 340 - 341 mols. of nucleotides, non-translation region comprising 1885 - 2551 mols. of nucleotides including region 1,149 and, etc.
                                                                                                                                                                                                                                                                                                                     Non-A Non-B hepatitis virus; structural region; cDNA to genomic RNA; detection; reagent; anti-Non-A Non-B hepatitis virus antibody; vaccine; antigen; epitope; diagnosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The sequence is that of NANB hepatitis virus HC-OM gene polypeptide CP-5-3. It may may be used in a system for detecting NANB hepatitis. This method is highly specific and sensitive, and can detect NANB hepatitis virus which could not be detected by conventional methods
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      specific; HCV; NANBH.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2 RGSRPSWGPTDPRHRSRNVG
                                                                                                                                                                                                                                                                                                                                                                                                          of Non-A Non-B hepatitis virus structural protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RGSRPTWGPTDPRHRSRNLG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      63; 73pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 90JP-00153401.
90JP-00304405.
                                                                                93JP-00126709
94JP-00032201
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                                                                                                                                             94EP-00108256
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                              entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     89.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 110; DB 2
Pred. No. 2e-08;
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      Sawada
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 2;
ŗ,
      Hosoda T,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 20;
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        Iwasaki
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local S
Matches 18
The sequences given in AAR25130-35 are peptides which have reactivity to the antibody against hepatitis C virus (HCV). They can be used on their own or as a mixture two different peptides. Using these peptides, HCV antibody positive patients can be detected and hepatitis caused by blood transfusion can be prevented
                                                                                                                                     New peptides acting as antigenic analogues of human hepatitis {\tt C} virus useful for detecting HCV antibody positive patients.
                                                                                                                                                                                          WPI; 1992-231947/28
                                                                                                                                                                                                                                                         19-OCT-1990;
                                                                                                                                                                                                                                                                                         19-OCT-1990;
                                                                                                                                                                                                                                                                                                                             02-JUN-1992.
                                                                                                                                                                                                                                                                                                                                                              JP04159298-A
                                                                                                                                                                                                                                                                                                                                                                                              Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                               Hepatitis C virus; blood transfusion
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    23-DEC-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAR25132 standard; protein; 20
                                                                                                     Claim 1; Page 1; 14pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 20
                                                                                                                                                                                                                          (OLYU ) OLYMPUS OPTICAL CO LTD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 3; Page 40; 59pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New non-A non-B hepatitis virus sub-type - used to develop prods. for detection, diagnosis, prevention and treatment of non-A non-B hepatitis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1995-015655/03
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         l Similarity 90.(
18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2 RGSRPSWGPTDPRHRSRNVG 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RGSRPTWGPTDPRHRSRNLG 20
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                                                                                                                                                                                                                                                           90JP-00282431.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     prevention and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 110; DB 2
Pred. No. 2e-08;
2; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0
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RESULT 12
AAR56608
ID AAR56
XX
AC AAR56
XX
DT 24-MA

AAR56608; AAR56608

standard;

protein;

20

A

24-MAR-1995

(first entry)

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RESULT 11
AAR38292
ID AAR38
XX AAR38
XX AAR38
XX NANAB
XX I2-UI
PR 08-NC
XX NAXU
PR 10-AI
XX NAXU
PR 12-UI
PR 08-NC
XX NAXU
PR N-PSI
XX NAXU
XX NAXU
XX I2-UI
PR N-PSI
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local S
Matches 18
                                                                                                   Query Match
Best Local Similarity
Matches 18; Conserv
                                                                                                                                                                                                                                                              The sequence is that of NANB hepatitis virus HC-OM gene polypeptide CP-5-
1. It may may be used in a system for detecting NANB hepatitis. This
method is highly specific and sensitive, and can detect NANB hepatitis
virus which could not be detected by conventional methods
                                                                                                                                                                                                             Sequence 20 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Antigen related to non-A and non-B hepatitis virus - comprises non translation region comprising 340 - 341 mols. of nucleotides, non-translation region comprising 1885 - 2551 mols. of nucleotides incregion 1,149 and, etc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           N-PSDB; AAQ43902.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              12-JUN-1990;
08-NOV-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Non-A, non-B; virus; specific; HCV; NANBH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 20 AA,
                                                                                                                                                                                                                                                                                                                                                                                                    Claim 28; Page 63; 73pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           10-APR-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        JP05091884-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NANB hepatitis virus HC-OM gene polypeptide CP-5-1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAR38292;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAR38292 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WPI; 1993-199637/25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           non-B hepatitis virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     16-APR-1993.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (NAKA/) NAKAMURA T.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      18;
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                                                RGSRPSWGPTDPRHRSRNVG 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RGSRPSWGPTDPRRRSRNLG 20
RGSRPSWGPTDPRRRSRNLG
                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              90JP-00153401.
90JP-00304405.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              91JP-00196175
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   protein;
                                                                                                                             90.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             90.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  polymerase chain reaction; detection; sensitive;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      20
                                                                                                                          Score 105;
Pred. No. 9.
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Pred. No. 9.9e-08;
20
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                                                                                                      Mismatches
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                                                                                                                             DB 2;
.9e-08;
                                                                                                                                                    Length 20;
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                                                                                                      Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
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                                                                                                      Gaps
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RESULT 13
AAU84604
ID AAU84
XX AAU84
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XX Savin
XX Savin
XX Savin
XX Savin
XX Savin
XX Savin
XX Hepa'
XX Hepa'
XX Savin
XX Fall
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Best Local S
Matches 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A T-cell stimulating peptide is claimed which is ca. 5-20 amino acids long and part of the core protein of HCV and is recognised by and stimulates T-cells. A CD8-positive T-cell stimulating peptide has the sequence given in AAR55606 (NP-9) or AAR56616 (9MA). A CD4- positive T-cell stimulating peptide has the sequence given in AAR56009 (NP-12) or AAR56614 (NP-17). Synthetic peptide mixts. (Mix A: NP-1 - NP-5, Mix B: NP-6 - NP-10, Mix C: NP-11 - NP14, MixD: NP-18) were used in experiments to evaluate activity. Peptides 9MA, 9CM and 9CA (AAR56616-18) are variants of peptide NP-9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ACA
                                                                                                                                                                                                                                       WO200190197-A1
                                                                                                                                                                                                                                                                                              Hepatitis C virus.
                                                                                                                                                                                                                                                                                                                                                                             Savine; vaccine; cancer; viral infection; HIV; hepatitis C virus; viral infection; human immunodeficieny virus; melanoma; bacterial infection; Salmonella; Legionella; parasitic infection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                HCV HepCla segment 7.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          08-MAY-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAU84604;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 20 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure; Fig 2; 14pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 1994-269451/33
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10-NOV-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        02-APR-1993;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Hepatitis C virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Hepatitis C virus; HCV; core protein; epitope; T-cell; immunity; CD8.
                                                            26-MAY-2000; 2000AU-00007761.
                                                                                                                                                                                29-NOV-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAU84604 standard; peptide; 30
                                                                                                                      25-MAY-2001; 2001WO-AU000622
                                                                                                                                                                                                                                                                                                                                                    Trypanosoma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (ASAH ) ASAHI KASEI KOGYO KK (INOM/) INOMAWARI M.
      (AUSU )
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 I cell epitope present in the core protein region of Hepatitis C virus (HCV) - used for activation of cellular immunity mechanisms.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        peptide NP-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 18;
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         UNIV AUSTRALIAN NAT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (first entry)
                                                                                                                                                                                                                                                                                                                                                 Toxoplasma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      92JP-00228965.
92JP-00299691.
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90.0%;
                                                                                                                                                                                                                                                                                                                                                       Giardia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 105; DB 2;
Pred. No. 9.9e-08;
1; Mismatches 1
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Sequence 30

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modulating immune responses preferably directed against a pathogen or a cancer, (e.g., cancers of the lung, breast, ovary, cervix, colon, head and neck, pancreas, prostate, stomach, bladder, kidney, bone liver, oesophagus, brain, testicle, uterus), as potentiating agents. Compositions comprising the polypeptide may be used in the treatment or prophylaxis against viral (such as infections caused by HIV (human immunodeficiency virus), hepatitis, influenza, Japanese encephalitis virus, Epstein-Barr virus and respiratory syncytial virus), bacterial (e.g., infections caused by Neisseria, Meningococcal, Haemophilus, Salmonella, Streptococcal, Legionella and Mycobacterium or parasitic (e.g., infections caused by Plasmodium, Schistosoma, Leishmania, Toxoplasma and Giardia) infections. The present sequence is
                                                                                                                                                                                                                                                                                                                                                                                                parent polypeptide to impede, abrogate or otherwise alter at least one function associated with the parent polypeptide and for inducing an immune response against a pathogen or cancer. Also included are a synthetic polynucleotide encoding and a computer system for designing the synthetic polypeptides. The synthetic polypeptides and polynucleotides are referred to as a Savine. The synthetic polypeptide is useful for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New synthetic polypeptides having several different segments of at least one parent polypeptide linked together differently compared to the linkage in the parent polypeptide, for inducing immune response against
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2002-147575/19.
N-PSDB; ABK36442.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The invention relates to a new synthetic polypeptide (I) comprising several different segments of at least one parent polypeptide linked together in a different relationship relative to their linkage in the
                                                                          Trypanosoma,
a peptide de
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 2; Fig 26; 364pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    pathogen or cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Thomson SA,
                                                                          derived
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Ramshaw
                                                                       from a parent protein used to construct a savine of
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RESULT 14
AAR31922
ID AAR31
XX AAR31
AC AAR31
XY DT 25-M
DT 03-M
XX Hepa
XX Hepa
XX Syni
XX Ep5
XX Ep5
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XX Ex6
XX Ep5
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XX (B
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Matches 18
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03-MAR-1993
                                05-APR-1991;
                                                   03-APR-1992;
                                                                       07-OCT-1992
                                                                                            EP507615-A1
                                                                                                              Synthetic.
                                                                                                                                 Hepatitis C
                                                                                                                                                     HCV peptide
                                                                                                                                                                                                        AAR31922;
                                                                                                                                                                                                                             AAR31922 standard;
                                                                                                                                                                                                                                                                               12
                                                                                                                                                                                                                                                                                                                      18;
                                                                                                                                                                                                                                                                                           1 PRGSRPSWGPTDPRHRSRN 19
                                                                                                                                                                                                                                                                                                                                Similarity
                                                                                                                                                                                                                                                                                PRGSRPSWGPTDPRRRSRN
                                                                                                                                   virus; vaccine; screening; diagnosis
                                                                                                                                                        BCH-443
                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                          (revised)
(first en
                               91US-00681701
                                                   92EP-00302975.
                                                                                                                                                                                                                            peptide;
                                                                                                                                                                                                                                                                                                                                85.4%;
94.7%;
                                                                                                                                                                           entry)
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Pred. No. 1.5e-07;
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(BIOC-) BIOCHEM PHARMA INC

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RESULT 15
AAM50830
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Best Local S
Matches 17
The present sequence is that of peptide MDL-8 comprising amino acid residues 103-120 of the hepatitis C virus (HCV) polyprotein core region (see AAM50816). A panel of overlapping peptides between positions 1 and 186 of the HCV core region was synthesised and tested for immunoreactivity with a commercially available human serum panel. All of the immunoreactivity was concentrated within the core region between amino acids 1 to 80. MDL-8 did not include a major immunoceactivity was concentrated virting the core region between The invention provides a highly immunoreactive mosaic antigen composition (MAC) comprising 2 or more immunoreactive peptides (see AAM50816-27), each of which is immobilised to a carrier. The unique combination of HCV core peptides in the MAC provides higher specificity and sensitivity for
                                                                                                                                                                                                                   New hepatitis C virus (HCV) peptides and mosaic antigen composition comprising the peptides, useful as immunoreagents for detecting HCV antibodies infection, or as an immunogen for stimulating production antibodies against HCV.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ACA
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01-MAY-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The peptide BCH-443 was synthesised based upon amino acids 101-127 of hepatitis C virus (HCV) envelope protein. It is useful as an active ingredient in vaccines against HCV and can be used for screening and diagnosing HCV infection. See also AAR31917-R31926. (Updated on 25-MAR-2003 to correct PN field.)
                                                                                                                                                                                        Example 1;
                                                                                                                                                                                                                                                                                               WPI; 2002-179696/23.
                                                                                                                                                                                                                                                                                                                               Chan HKW,
                                                                                                                                                                                                                                                                                                                                                          (MEDM-) MEDMIRA INC.
                                                                                                                                                                                                                                                                                                                                                                                     07-JUL-2000; 2000CA-02311022
                                                                                                                                                                                                                                                                                                                                                                                                                     06-JUL-2001; 2001WO-CA000988
                                                                                                                                                                                                                                                                                                                                                                                                                                                  17-JAN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO200204484-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Hepatitis C virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         HCV; MDL-8; infection; diagnosis; vaccine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAM50830;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAM50830 standard; peptide; 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Hepatitis C virus peptide(s) and their mixts. - u vaccinating against hepatitis C virus infections.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 1; Page 27; 34pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 1992-333826/41.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Lacroix
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         core
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      region peptide MDL-8
                                                                                                                                                                                                                                                                                                                             Theolis R;
                                                                                                                                                                                      Page 33; 59pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (revised)
(first entry)
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Pred. No. 6.6e-07;
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    used for diagnosing

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                                                                                                                    Query Match
Best Local S
Matches 16
                                                                                                                                             detection of human antibodies specificity to HCV in rapid HCV diagnostic applications. An in vitro diagnostic method for detecting anti-HCV antibodies in a test sample and a diagnostic test kit are provided which use the MAC as an immunoreagent. The HCV MAC may also potentially be used as an immunogen in vaccine compositions. (Updated on 07-AUG-2003 to correct OS field.)
                                                                                                                    Sequence 18 AA;
                                                                         Local Similarity
                                                        16;
                            4 SRPSWGPTDPRHRSRNVG
SRPSWGPTDPRRRSRNLG
                                                         Conservative
                                                                       76.4%;
88.9%;
                                                     Score 94; DB 5; Len;
; Pred. No. 3.1e-06;
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 18
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Maximum DB
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Perfect score:
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   Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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                      US-10-13-007-5
PCT-US92-07865-15
US-08-995-0548-3
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PCT-US92-07813-8
US-08-974-690C-9
US-08-447-4918-301
US-08-444-818-301
US-08-444-818-300
US-08-444-818-300
US-08-38-34-818-300
US-09-308-3458-31
US-09-314-268-110
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US-09-318-348-34
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62.689 Million cell updates/sec
                                         Sequence 5, Appli
Sequence 3, Appli
Sequence 3, Appli
Sequence 3, Appli
Sequence 8, Appli
Sequence 8, Appli
Sequence 8, Appli
Sequence 22, Appli
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Sequence 23, Appli
Sequence 30, Appli
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US-08-398-028B-6	US-08-460-343B-6	US-09-314-268-109	US-09-504-262D-68	US-09-184-658-68	US-08-467-344A-306	US-08-488-446-306	US-08-469-260A-306	US-09-076-721-47	US-08-244-701B-47	US-08-974-690C-54	US-08-635-886C-54	US-08-951-034B-14	US-08-676-279-12	US-08-676-279-6	US-09-308-345A-33	US-09-308-345A-32	US-09-101-059-11
Sequence 6, Appi	Sequence 6, App	Sequence 109, Ap	Sequence 68, App	Sequence 68, App	•	Sequence 306, App	-	Sequence 47, Appl	•	•	Seguence 54, App	Sequence 14, App	Sequence 12, App	Sequence 6, Appl	Sequence 33, App	Sequence 32, App	Sequence 11, App

#### ALIGNMENTS

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APPLICANT: Ohue, Chiharu
APPLICANT: Ohue, Chiharu
APPLICANT: Iida, Kumiko
APPLICANT: 1ida, Kumiko
APPLICANT: Yagi, Shintaro
APPLICANT: Yagi, Shintaro
APPLICANT: Yagi, Shintaro
CURRENT APPLICATION NUMBER: US/10/133,007
CURRENT FILING DATE: 2002-04-26
PRIOR APPLICATION NUMBER: US/99/509,449
PRIOR APPLICATION NUMBER: UP 10-216094
PRIOR APPLICATION NUMBER: JP 10-216094
PRIOR APPLICATION NUMBER: JP 10-216094
PRIOR FILING DATE: 1998-07-30
PRIOR APPLICATION NUMBER: JP 10-216094
PRIOR FILING DATE: 1999-07-30
PRIOR FILING DATE: 1999-07-30
PRIOR FILING DATE: 1999-07-30
INMBER OF SEQ ID NOS: 11
SOFTWARE: Patentin version 3.1
SQC ID NO 5
LENGTH: 21
TYPE: PRT
ORGANISM: Artificial Sequence
PEATURE:
COTHER TNFORMATION: Fused polypeptide including Hepatitis C virus
                     Sequence 15, Application PC/TUS9207865
GENERAL INFORMATION:
APPLICANT: Dreesman, Gordan R.
APPLICANT: Burk, Kenneth H.
APPLICANT: Burk, Kenneth H.
APPLICANT: Pauletti, Daniel
TITLE OF INVENTION: Peptide-Based Hej
TITLE OF INVENTION: Immunoassays
NUMBER OF SEQUENCES: 25
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                 RESULT 2
PCT-US92-07865-15
                                                                                                                                                                                                                                                                                                                                                     밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; OTHER INFORMATION: Fused polypeptide including Hepatitis C virus US-10-133-007-5
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US-10-133-007-5
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Best Local S
Matches 21
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Patent No. 6623921
  ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  h 100.0%; Score 123; DB 4; Similarity 100.0%; Pred. No. 2.6e-11; Conservative 0; Mismatches 0;
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Law Offices of Peter Dehlinger
                                                                                               Hepatitis C
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; FRAGMENT TYPE: internal
; ORIGINAL SOURCE:
; ORGANISM: Hepatitis C Virus
; INDIVIDUAL ISOLATE: DP10, an
; INDIVIDUAL ISOLATE: polyprot
PCT-US92-07865-15
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                                                PRIOR APPLICATION NUMBER: US 08/707,355
PRIOR FILING DATE: 1996-09-04
PRIOR APPLICATION NUMBER: US 08/507,740
PRIOR HILING DATE: 1995-07-26
PRIOR APPLICATION NUMBER: US 08/373,920
PRIOR APPLICATION NUMBER: US 08/373,920
PRIOR FILING DATE: 1995-01-17
PRIOR APPLICATION NUMBER: US 08/183,207
PRIOR APPLICATION NUMBER: US 07/760,292
PRIOR FILING DATE: 1994-01-18
PRIOR APPLICATION NUMBER: US 07/760,292
PRIOR FILING DATE: 1991-09-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEPHONE: (415) 324-0860
TELEPAX: (415) 324-0860
INFORMATION FOR SEQ ID NO: 15:
SEQUENCE CHARACTERISTICS:
LENGTH: 24 amino acids
TYPE: ANINO ACID
                                                                                                                                                                                                                                                                                                                                                                                      Sequence 3, Application US/08905054B
Patent No. 6596476
GENERAL INFORMATION:
APPLICANT: Abbott Laboratories
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Matches
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                                                                                                                                                                                                                                                               FILE REFERENCE: 4767.US.C7
CURRENT APPLICATION NUMBER: US/08/905,054B
CURRENT FILING DATE: 1997-08-01
                                                                                                                                                                                                                                                                                                                             APPLICANT: Lesniewski, Richard R. APPLICANT: Leung, Tat K. TITLE OF INVENTION: HEPATITIS C ASSAY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TOPOLOGY: linear MOLECULE TYPE: peptide HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Fabian, Gary R.
REGISTRATION NUMBER: 33,875
REFERENCE/DOCKET NUMBER: 1600-0086.41
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: I
FILING DATE: 19920910
CLASSIFICATION:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ANTI-SENSE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILING DATE: 16-SEP-1991
FILING DATE: 12-FEB-1992
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STREET: 350 Cam
CITY: Palo Alto
STATE: CA
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               APPLICATION NUMBER: US 07/610,180 FILING DATE: 1990-11-07
APPLICATION NUMBER: US 07/456,162
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              l Similarity
19; Conserv
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NO: 15:
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                                                                                                                                                                                                                                                                                                                                                                     Richard R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DP10, amino acids 98-121 of the HCV polyprotein
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Pred. No. 1.1e-09;
1; Mismatches 1
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; ORGANISM: Hepatitis C Virus US-08-905-054B-3
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PCT-US92-07813-3
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                                                              RESULT 5
US-07-681-701-10
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Best Local S
Matches 19
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NUMBER OF SEQ ID NOS: 30
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 3
Sequence 10, Application US/07681701
Patent No. 5574132
GENERAL INFORMATION:
APPLICANT: Lacroix, Martial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 3, Application GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                               TELEFAX: 708-937-9556
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 28
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                       NAME: POREMBSKIP, PRISCILLA REGISTRATION NUMBER: 33,207 REFERENCE/DOCKET NUMBER: 476 TELECOMMUNICATION INFORMATION: TELEPHONE: 708-937-6365
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: PCT/I
FILING DATE: 19920916
CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: LESNIEWSKI, RICHARD R.
APPLICANT: LEUNG, TAT K.
TITLE OF INVENTION: HEPATITIS C ASSAY
NUMBER OF SEQUENCES: 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                   MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                TYPE:
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STREET: ONE ABBOTT PARK ROAD
CITY: ABBOTT PARK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY: U.S.A.
ZIP: 60064-3500
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LEUNG, TAT K.
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pred. No. 1.3e-09;
1; Mismatches 1
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Pred. No. 1.3e-09;
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                                                                                                                                                                                                                        Length 28;
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FILE REFERENCE: 2752-18

CURRENT APPLICATION NUMBER: US/08/635,886C

CURRENT FILING DATE: 1996-04-25

PRIOR APPLICATION NUMBER: PCT/EP94/03555

PRIOR FILING DATE: 1994-10-28

PRIOR APPLICATION NUMBER: EP 9402718.6

PRIOR FILING DATE: 1993-11-04

NUMBER OF SEQ ID NOS: 286

SOFTWARE: PatentIn version 3.1

SEQ ID NO 8

LENGTH: 20
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                                                                                     US-08-635-886C-8
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MOLECULE TYPE: peptide
US-07-681-701-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEX: 14-8367
INFORMATION FOR SEQ ID NO: 1
SEQUENCE CHARACTERISTICS:
LENGTH: 27 amino acids
TYPE: AMINO ACID
TYPE: AMINO ACID
Query Match
Best Local Similarity 94.
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 8, Application US/08635886C
Patent No. 6555114
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 81.3%;
Best Local Similarity 85.0%;
Matches 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: LEROUX-ROELS, Geert
APPLICANT: DELEYS, Robert
APPLICANT: MARGENS, Geert
TITLE OF INVENTION: IMMUNODOMINANT HUMAN T CELL EPITOPES OF HEPATITIS C
TITLE OF INVENTION: VIRUS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME: Haley Jr., James F.
REGISTRATION NUMBER: 27,794
REFERENCE/DOCKET NUMBER: IAF-10
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 715-0742
TELEPAX: (212) 715-0673
TELEX: 14-8367
                                                                                                TYPE: PRT
ORGANISM: hepatitis C virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TITLE OF INVENTION: PEPTIDES
TITLE OF INVENTION:
NUMBER OF SEQUENCES: 17
CORRESPONDENCE ADDRESS:
ADDRESSEE: FISH & NEAVE
STREET: 875 THird Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION DATA:
APPLICATION 1930405
FILING DATE: 19310405
CLASSIFICATION: 530
CLASSIFICATION: 530
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RGSRPSWGPNDPRRRSRNLG 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ss: single
linear
                  76.4%;
94.1%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   10:
Score 94; DB 4;
Pred. No. 3.5e-07;
0; Mismatches 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 100; DB 1;
Pred. No. 6.7e-08;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
                                        Length 20;
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APPLICANT: DELEYS, ROBERT

APPLICANT: MARRITENS, GeerE

TITLE OF INVENTION: IMMUNODOMINANT HUMAN T CELL EPITOPES OF HEPATITIS C

TITLE OF INVENTION: VIRUS

FILE REFERENCE: 2551-94

CURRENT APPLICATION NUMBER: US/08/974,690C

CURRENT APPLICATION NUMBER: PCT/EP94/0355

PRIOR APPLICATION NUMBER: PCT/EP94/0355

PRIOR APPLICATION NUMBER: EP 93402718.6

PRIOR APPLICATION NUMBER: EP 93402718.6

PRIOR FILING DATE: 1993-11-04

NUMBER OF SEQ ID NOS: 286

SOFTWARE: Patentin version 3.1

SEQ ID NO 8

LENGTH: 20

TYZEE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 8
US-08-604-365-22
; Sequence 22, Application US/08604365
; Patent No. 6183949
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US-08-974-690C-8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Matches
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 5.25 inch, 36
COMPUTER: IBM PS/2
OPERATING SYSTEM: PC-DOS
OPERATING SYSTEM: PC-DOS
SOTWARE: Wordperfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/604,365
FILING DATE:
CLASSIFICATION 1435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/977,398
FILING DATE: 11-MARCH-1993
APPLICATION NUMBER: PCT/EP92/01468
FILING DATE: 30-UNNE-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
APPLICANT: Seidel, Christoph; Ehrich-Weinreich,
APPLICANT: Gertraud; Bayer, Hubert; Wienhues, Ursula; Ju
APPLICANT: G nther-Gerhard; Ihlenfieldt, Hans Georg
TITLE OF INVENTION: HCV Peptide Antigens and Methods for
TITLE OF INVENTION: the Determination of HCV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: LEROUX-ROELS, Geert
                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES: 3
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                            COUNTRY: US
ZIP: 10022
                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE: Felfe & Lynch
STREET: 805 Third Avenue
CITY: New York City
STATE: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          16;
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                                                                                                                                                                                                                                                                                                                                   USA
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                                                                                                                                                                                                                                           E: Diskette, 5.25 inch, 360 kb storage IBM PS/2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            76.4%;
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Pred. No. 3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION UNMER: DE 41 41 304.0
PRILING DATE: 14-DEC-1991
PRIOR APPLICATION DATA:
DE 42 09 216.9
PRILING DATE: 21-MARCH-1992
ATTORNEY/AGENT INFORMATION:
NAME: TSai, Christine H.
REGISTRATION NUMBER: 34,266
REFERENCE/DOCKET NUMBER: BOER 1010
REGISTRATION INFORMATION:
TELEPHONE: (212) 688-9200
TELEPAX: (212) 688-9200
TELEPAX: (212) 838-3884
INFORMATION FOR SEQ ID NO: 22:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
TYPE: amino acids
TYPE: amino acids
TYPE: amino acids
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Best Local Similarity
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                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 5.25 inch, 360 kb storage
COMPUTER: IBM PS/2
COMPUTER: IBM PS/2
OPERATING SYSTEM: PC-DOS
SOFTWARE: WOrdperfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/689,678
FILING DATE: 13-Oct-2000
CLASSIFICATION: cUnknown>
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
ADDRESSEE: Felfe & Lynch
STREET: 805 Third Avenue
CITY: New York City
STATE: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Seidel, Christoph; Ehrich-Weinreich,
Gertraud; Bayer, Hubert; Wienhues, Ursula; Jun
Gunther-Gerhard; Inhenfeldt, Hans Georg
TITLE OF INVENTION: HCV Peptide Antigens and Methods for
the Determination of HCV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES: 32
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      14;
              APPLICATION NUMBER: US/08/604,365
FILLING DATE: «Unknown»
APPLICATION NUMBER: 07/977,398
FILLING DATE: 11-MARCH-1993
APPLICATION NUMBER: PCT/EP92/01468
FILLING DATE: 30.-JUNE-1992
APPLICATION NUMBER: DE 41 22 160.5
FILLING DATE: 04-JULY-1991
APPLICATION NUMBER: DE 41 41 304.0
FILLING DATE: 14-DEC-1991
APPLICATION NUMBER: DE 42 09 216.9
FILLING DATE: 21-MARCH-1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRGSRPSWGPTDPR 14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Application US/09689678
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 69.1%; Score 85; ilarity 100.0%; Pred. No. Conservative 0; Mismatc
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04-JULY-1991
TE: 21-MARCH-1992
INFORMATION:
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US-08-473-475A-25; Sequence 25, Ap; Patent No. 5843
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                                                                                                                                  ; MOLECULE TYPE:
US-08-473-475A-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 69.1%; Sometime 100.0%; East Local Similarity 100.0%; East Local Similarity 0; Matches 14; Conservative 0;
Query Match
Best Local Similarity
Matches 10; Conserv
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                                                                                                                                                                                                                                                                                          TELEFAX: 708-937-2623
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 31 amino acids
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CURRENT APPLICATION NUMBER: US/08/473 ATTELING DATE:

CLASSITETT:

CL
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT:
APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                REFERENCE/DOCKET NUMBER: 55
TELECOMMUNICATION INFORMATION:
TELEPHONE: 708-937-6365
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: POREMBSKI, PRISCIL:
REGISTRATION NUMBER: 332
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TITLE OF INVENTION: HEPATITIS GB VIRUS SYNTHETIC PEPTIDES TITLE OF INVENTION: AND USES THEREOF NUMBER OF SEQUENCES: 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STATE: IL COUNTRY: US ZIP: 60064
                                                                                                                                                                                             LENGTH: 31 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                           TELEPHONE: 708-37-2623
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STREET:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADDRESSEE:
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MOLECULE TYPE: peptide
SEQUENCE DESCRIPTION: SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: Tsai, Christine H.
REGISTRATION NUMBER: 34,266
REFERENCE/DOCKET NUMBER: BOER 1010
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 688-9200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 PRGSRPSWGPTDPR 14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            E: ABBOTT LABORATORIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DAWSON, GEORGE
BRIDON, DOMINIQUE P.
SCHROEDER-POLIAK, PAMELLA
KNIGGE, MARK F.
JAFFE, KZEVE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      USA
      Conservative
                                                                                                                                                              protein
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                                  52.0%;
71.4%;
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100.0%; Pred. No. 5.1e-06;
Live 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    838-3884
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   Score 64; DB 2; pred. No. 0.011; 2; Mismatches
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WGPTDPRHRSRNVG

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                                                                                                                              ; TYPE: PRT; ORGANISM: hepatitis C virus US-08-974-690C-9
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US-08-974-690C-9
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                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
APPLICANT: LEROUX-ROELS, Geert
APPLICANT: DELEYS, Robert
APPLICANT: MAERIENS, Geert
TITLE OF INVENTION: INMUNODOMINANT HUMAN T CELL EPITOPES OF HEPATITIS
TITLE OF INVENTION: VIRUS
                                                                                                                                                                              SOFTWARE: PatentIn version 3.1
SEQ ID NO 9
LENGTH: 20
                                                              Matches
                                                                            Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 9, Application US/08974690C Patent No. 6613333
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQ ID NOS: 286
SOFTWARE: PatentIn version 3.1
SEQ ID NO 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
                                                                                                                                                                                                                                                                              FILE REFERENCE: 2551-94
CURRENT APPLICATION NUMBER: US/08/974,690C
CURRENT FILING DATE: 1997-11-19
PRIOR APPLICATION NUMBER: PCT/EP94/03555
PRIOR FILING DATE: 1994-10-28
                                                                                                                                                                                                                               PRIOR APPLICATION NUMBER: EP 93402718.6
PRIOR FILING DATE: 1993-11-04
NUMBER OF SEQ ID NOS: 286
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: LEROUX-ROBLS, Geert
APPLICANT: DELEYS, Robert
APPLICANT: MAERTENS, Geert
TITLE OF INVENTION: IMMUNODOMINANT HUMAN T CELL EPITOPES OF HEPATITIS
TITLE OF INVENTION: VIRUS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION NUMBER: PCT/EP94/03555
PRIOR FILING DATE: 1994-10-28
PRIOR APPLICATION NUMBER: EP 93402718.6
PRIOR FILING DATE: 1993-11-04
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CURRENT APPLICATION NUMBER: US/08/635,886C CURRENT FILING DATE: 1996-04-25
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TYPE: PRT
ORGANISM: hepatitis C virus
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                            10 PTDPRHRSRNVG 21
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                                                              Conservative
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                                                                            46.3%;
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                                                                            Score 57;
Pred. No.

    Mismatches

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Pred. No. 0
                                                                          DB 4;
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                                                                                          Length 20;
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APPLICANT: ADVAGI, Katsumi
APPLICANT: Ohue, Chiharu
APPLICANT: Ohue, Chiharu
APPLICANT: 11da, Kumiko
APPLICANT: Yagi, Shintaro
TITLE OF INVENTION: METHOD FOR MEASUREMENT OF HEPATITIS C VIR
FILE REFERENCE: 594.352USWO
CURRENT APPLICATION NUMBER: US/10/133,007
CURRENT APPLICATION NUMBER: US/09/509,449
PRIOR APPLICATION NUMBER: US/09/509,449
PRIOR FILING DATE: 2000-03-28
PRIOR APPLICATION NUMBER: DT 10-216094
PRIOR FILING DATE: 1998-07-30
PRIOR FILING DATE: 1998-07-30
PRIOR FILING DATE: 1999-07-30
NUMBER OF SEQ ID NOS: 11
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Best Local Similarity
Matches 10; Conserve
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                                 COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION UMBER: US/08/444,818
FILING DATE:
CLASSIFICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: US/08/403,590
FILING DATE: 14-MAR-1995
ATTORNEY/AGENT INFORMATION:
NAME: Harbin, Alisa A.
REGISTRATION NUMBER: 33,895
                                                                                                                                                                                                                                                                                                                                                                                                                                               Patent No. 6150087
GENERAL INFORMATION:
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Chien, David Y.
APPLICANT: Rutter, William J.
TITLE OF INVENTION: NAMBY Diagnostics
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: PatentIn version 3.1
SEQ ID NO 6
LENGTH: 20
TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FEATURE:
OTHER INFORMATION: Fused polypeptide including Hepatitis C
REGISTRATION NUMBER: 33,895
REFERENCE/DOCKET NUMBER: 01
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
ADDRESSEE: Chiron Corporation
STREET: 4560 Horton Street
                                                                                                                                                                                                                                                                                                                                                                        STREET: 4560 Hort
CITY: Emeryville
                                                                                                                                                                                                                                                                                                                                        COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            6623
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ilarity 100.0%;
Conservative
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COMPUTER READABLE FORM:

COMPUTER READABLE FORM:

COMPUTER: IBM PC COMPATIBLE

COMPUTER: IBM PC COMPATIBLE

COMPUTER: IBM PC COMPATIBLE

COMPUTER: IBM PC COMPATIBLE

SOFTWARE: PATENTIN PC-DOS/MS-DOS

SOFTWARE: PATENTIN PC-DOS/MS-DOS

CUURENT APPLICATION DATA:

APPLICATION NUMBER: US/08/444,818

FILING DATE: 14-WAR-1995

APPLICATION NUMBER: US/08/403,590

APPLICATION NUMBER: US/08/403,590

APPLICATION NUMBER: US/08/403,590

APPLICATION NUMBER: US/08/403,590

APPLICATION NUMBER: 33,895

REGISTRATION NUMBER: 0110.002

TELEPHONE: (508)359-3876

TELEPHONE: (508)359-3876

TELEPHONE: (508)359-3876

INFORMATION FOR SEQ ID NO: 299:

SEQUENCE CHARACTERISTICS:

LENGTH: 8 amino acids

TYPE: amino acids

TYPE: amino acids

STRANDEDNESS: single

TOPOLOGY: linear

MOLECULE TYPE: peptide

US-08-444-818-299

39.8%; Score 49; DB 3; Le

39.8%; Score 49; DB 3; Le
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 299, Application US/08444818

Patent No. 615087

GENERAL INFORMATION:
APPLICANT: Chien, David Y.
APPLICANT: Rutter, William J.
TITLE OF INVENTION: NAMBY Diagnostics and Vaccines NUMBER OF SEQUENCES: 777

CORRESPONDENCE ADDRESS:
ADDRESSEE: Chiron Corporation
STREET: 4560 Horton Street
CTIY: Emeryville
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MOLECULE TYPE: peptide US-08-444-818-301
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US-08-444-818-299
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SEQUENCE CHARACTERISTICS:
LENGTH: 8 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
Search completed: June 29, 2004, 12:09:59 Job time: 17.2941 secs
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                                                                                                                                                             Query Match
Best Local S
Matches 8
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ZIP: 94608-2916
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                                                                                                                     PRGSRPSW 8
                                                                                   PRGSRPSW 8
                                                                                                                                                           39.8%; Score 49; DB 3; ilarity 100.0%; Pred. No. 3e+05; Conservative 0; Mismatches
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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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Perfect score:
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Gapop 10.0 , Gapext 0.5
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137.314 Million cell updates/sec
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123
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/ Ggn2_6/ptodata/1/pubpaa/USO7_PUBCOMB.pep:*
/ Ggn2_6/ptodata/1/pubpaa/USO7_NEW_PUB.pep:*
/ Ggn2_6/ptodata/1/pubpaa/USO6_NEW_PUB.pep:*
/ Ggn2_6/ptodata/1/pubpaa/USO6_PUBCOMB.pep:*
/ Ggn2_6/ptodata/1/pubpaa/USO7_NEW_PUB.pep:*
/ Ggn2_6/ptodata/1/pubpaa/USO8_PUBCOMB.pep:*
/ Ggn2_6/ptodata/1/pubpaa/USO8_PUBCOMB.pep:*
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/ Ggn2_6/ptodata/1/pubpaa/USO9E_PUBCOMB.pep:*
/ Ggn2_6/ptodata/1/pubpaa/USO9E_PUBCOMB.pep:*
/ Ggn2_6/ptodata/1/pubpaa/USO9E_PUBCOMB.pep:*
/ Ggn2_6/ptodata/1/pubpaa/USO9E_PUBCOMB.pep:*
/ Ggn2_6/ptodata/1/pubpaa/USO9E_PUBCOMB.pep:*
/ Ggn2_6/ptodata/1/pubpaa/USO9E_PUBCOMB.pep:*
/ Ggn2_6/ptodata/1/pubpaa/USO0E_PUBCOMB.pep:*
/ Ggn2_6/ptodata/1/pubpaa/USO0E_PUBCOMB.pep:*
/ Ggn2_6/ptodata/1/pubpaa/USO0E_PUBCOMB.pep:*
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/ Ggn2_6/ptodata/1/pubpaa/USO0E_PUBCOMB.pep:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

# SUMMARIES

Result

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49	54	57	1 or	, do	67	67	78	78	80	90	94	105	112	112	Score
39.8	43.9	46.3	52.8		٠		63.4	63.4	69.1	٠		85.4	91.1	ŗ	Match
v	18	20	15	15	15	15	25	25	15	30	20	30	25	25	Match Length
ω	15	12	14	14	14	14	14	10	14	12	12	12	14	10	DB
US-08-344-824-83	US-10-350-719-180	US-10-651-165-9	US-10-268-569-12	US-10-268-561-12	US-10-268-569-11	US-10-268-561-11	US-10-173-480-57	US-09-891-983A-30	US-10-371-540-22	US-10-296-734-422	US-10-651-165-8	US-10-296-734-420	US-10-173-480-56	US-09-891-983A-29	ID
Sequence 83, Appl	Sequence 180, App	9,	12,	12,	11,	11,	57,	30,	22,	422	8	Ф 4.	e 56,	Sequence 29, Appl	Description

RESULT 2 US-10-173-480-56

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38	38	38 8	38	38	œ	ω.	38.5	39	39	39	39	9	39.5	9	40	40	40	40	0	40.5	0	42	44		44	44	44.5	49	49
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-09-305-736	-09-974-879-56	0-084-813-34	-10-084-813-34	0-084-813-34	19-	-10-008-524A-1	9-864-761-3745	-880-578-36	LO-194-441A	10-440-390-2	3-344-824-84	-10-350-719-10	-10-008-524A-1	0-029-386-314	-10-144-929-25	-10-144-929-25	00-303A-4	-10-228-806-61	25-302	9-925-302-68	9-775-964-2	09-836-392-3	α	-10-023-282-63	-10-004-860-6	-09-933-767-63	9-38	-10-173-480-5	US-09-891-983A-28
Sequence 569, App	emience 564	equence 349.	equence 348,	equence 347,	e 107,	Sequence 107,	nce 3745	quence 36, Appl	equence 83,	ence 29, Āpp	quence 84, Appl	equence 106, Ap	nce 106, Ap	equence 31	e 252, Ap	equence 25	e 48	e 61	е 6	681	ce 28	30,	e 28430	equence 630,	equence 630	equence 630,	equence 33805,	e 55, App	8, Ap

#### ALIGNMENTS

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APPLICANT: Leary, Thomas P.
APPLICANT: Desai, Suresh
APPLICANT: Stewart, James L.
TITLE OF INVENTION: Methods For The simultaneous Detection
TITLE OF INVENTION: Of HCV Antigens And HCV Antibodies
FILE REFERENCE: 6821.US.01
CURRENT APPLICATION NUMBER: US/09/891,983A
CURRENT FILING DATE: 2001-06-26
NUMBER OF SEQ ID NOS: 36
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 29
LENGTH: 25
TYPE: PAT
ORGANISM: Hepatitis C Virus
US-09-891-983A-29
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US-09-891-983A-29
                                                                     Query Match
Best Local Similarity
Matches 19; Conserv
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APPLICANT: Abbott Laboratories
APPLICANT: Shah, Dinesh O.
APPLICANT: Dawson, George A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 29, Application US/09891983A Publication No. US20030108858A1
                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT:
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                          1 PRGSRPSWGPTDPRHRSRNVG 21
IJ
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Muerhoff, A. Scott
Jiang, Lily
Pohin A.
PRGSRPSWGPTDPRRRSRNLG 25
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                                                                     Conservative
                                                                                    91.1%;
                                                                   1;
                                                                                  Score 112; DB 10;
Pred. No. 1.7e-07;
                                                                     Mismatches
                                                                                                   Length 25;
                                                                   Indels
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                                                                 Gaps
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; Sequence 8, Application US/10651165
                  RESULT 4
US-10-651-165-8
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US-10-296-734-420
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                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION NUMBER: US/10/296,734
CURRENT FILING DATE: 2003-08-04
PRIOR APPLICATION NUMBER: AU PQ7761/00
PRIOR FILING DATE: 2000-05-26
NUMBER OF SEQ ID NOS: 1507
SOFTWARE: PatentIn version 3.2
SEQ ID NO 420
SEQ ID NO 420
                                                                                                                                                                                          Matches
                                                                                                                                                                                                            Query Match
Best Local (
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LENGTH: 25
TYPE: PRI
TYPE: PRI
ORGANISM: Artificial Sequence
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Best Local Similarity
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APPLICANT:
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TITLE OF INVENTION: METHODS FOR THE SIMULTANEOUS DETECTION

TITLE OF INVENTION: OF HCV ANTIGENS AND HCV ANTIBODIES

FILE REFERENCE: 6821.US.P1

CURRENT APPLICATION NUMBER: US/10/173,480

CURRENT FILING DATE: 2002-06-17

PRIOR APPLICATION NUMBER: 09/891,983

PRIOR APPLICATION NUMBER: 09/891,983

PRIOR PILING DATE: 2001-06-26

NUMBER OF SEQ ID NOS: 63

NUMBER OF SEQ ID NOS: 63
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APPLICANT: Shah, Dinesh O.
APPLICANT: Dawson, Greoge A.
APPLICANT: Muerhoff, A. Scott
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Ramshaw, Ian A
                                                                                                                                                                                                                                                                                                                       LENGTH: 30
TYPE: PRT
ORGANISM: Artificial
                                                                                                                                                                                                                                                                                                                     FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           OTHER INFORMATION: HCV-Core derived peptides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FEATURE:
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                                                                                                                                             1 PRGSRPSWGPTDPRHRSRN 19
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                                                                                                                                                                                                            Similarity
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                                                                                                      PRGSRPSWGPTDPRRRSRN 30
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Leary, Thomas P.
Desai, Suresh
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                                                                                                                                                                                        Conservative
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90.5%;
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94.7%;
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Pred. No. 1.6e-06;
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                                                                                                                                                                                        Mismatches
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                                                                                                                                                                                                                              Length 30;
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APPLICANT: Thompson, Scott A
APPLICANT: Ramshaw, Ian A
TITLE OF INVENTION: Synthetic molecules and uses therefor
FILE REFERENCE: Savine
CURRENT APPLICATION NUMBER: US/10/296,734
CURRENT FILING DATE: 2003-08-04
PRIOR APPLICATION NUMBER: AU PQ7761/00
PRIOR APPLICATION NUMBER: AU PQ7761/00
PRIOR FILING DATE: 2000-05-26
NUMBER OF SEQ ID NOS: 1507
SOFTWARE: Patentin version 3.2
SEQ ID NO 422
LENGTH: 30
TYPE: PRT
ORGANISM: Artificial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION NUMBER: US/08/974,690C
PRIOR FILING DATE: 1997-11-19
PRIOR APPLICATION NUMBER: PCT/EP94/0355
PRIOR FILING DATE: 1994-10-28
PRIOR APPLICATION NUMBER: EP 93402718.6
PRIOR FILING DATE: 1993-11-04
NUMBER OF SEQ ID NOS: 286
SOFTMARE: PATENTIAN VERSION 3.1
SEQ ID NO 8
LENGTH: 20
; Sequence 22, Application US/10371540
; Publication No. US20030198644A1
; GENERAL INFORMATION:
; APPLICANT: Roche Diagnostics GmbH
; TITLE OF INVENTION: HCV PEPTIDE ANTIGENS AND METHODS FOR THE DETERMINATION OF HCV
; FILE REFERENCE: 9793/129
                                                                                                                            RESULT 6
US-10-371-540-22
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; Publication No. US20
; GENERAL INFORMATION:
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US-10-296-734-422
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                                                                                                                                                                                                                                                                                                         Matches
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GENERAL INFORMATION:
APPLICANT: LEROUX-ROELS, Geert
APPLICANT: MAERIENS, Geert
APPLICANT: MAERIENS, Geert
TITLE OF INVENTION: INMUNDOMINANT HUMAN T CELL EPITOPES OF HEPATITIS C
TITLE OF INVENTION: VIRUS
                                                                                                                                                                                                                                                                                                                           Query Match
Best Local :
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Best Local Similarity
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CURRENT APPLICATION NUMBER: US/10/651,165
CURRENT FILING DATE: 2003-09-02
                                                                                                                                                                                                                                                                                                                                                                                                                                   FEATURE:
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                                                                                                                                                                                                                                                                                                      Conservative
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94.1%;
                                                                                                                                                                                                                                                                                                                           73.2%;
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Pred. No. 2.8e-05;
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RESULT 8
US-10-173-480-57
/ Sequence 57, Application US/10173480
/ Publication No. US20030152948A1
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                                                                                                                                                                                                                                                                                                                                         APPLICANT: Stewart, James L.
TITLE OF INVENTION: Methods For The simultaneous Detection
TITLE OF INVENTION: Methods For The simultaneous Detection
FILE REFERENCE: 6821.US.01
CURRENT APPLICATION NUMBER: US/09/891,983A
CURRENT FILING DATE: 2001-06-26
NUMBER OF SEQ ID NOS: 36
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 30
LENGTH: 25
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-09-891-983A-30
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CURRENT FILING DATE: 2003-02-21
PRIOR APPLICATION NUMBER: 09/689,678
PRIOR FILING DATE: 2000-10-13
PRIOR APPLICATION NUMBER: 08/604,365
PRIOR APPLICATION NUMBER: 04/14 1 304.1
PRIOR APPLICATION NUMBER: P 41 41 304.1
PRIOR APPLICATION NUMBER: P 42 09 215.9
PRIOR APPLICATION NUMBER: P 47 09 215.9
PRIOR APPLICATION NUMBER: P 47 09 215.9
PRIOR APPLICATION NUMBER: P 47 22 160.5
PRIOR APPLICATION NUMBER: P 41 22 160.5
PRIOR FILING DATE: 1992-06-30
PRIOR FILING DATE: 1991-07-04
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Best Local S
Matches 13
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SOFTWARE: PatentIn version 3.1
SEQ ID NO 22
LENGTH: 15
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APPLICANT: Shah, Dinesh O.
APPLICANT: Dawson, George A.
APPLICANT: Dawson, George A.
APPLICANT: Muerhoff, A. Scott
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TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                ORGANISM: Hepatitis C Virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FEATURE: OTHER INFORMATION: peptide antigen for anti-HCV antibodies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local
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                                                                                                                                                                                7
                                                                                                                                                                                                                      l Similarity
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Similarity 100.0%; Pred. No. 0.00031
14; Conservative 0; Mismatches (
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                                                                                                                                    SWGPTDPRRRSRNLG
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Leary, Thomas P.
Desai, Suresh
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No. US20030108858A1
                                                                                                                                                                                                                                         63.4%;
                                                                                                                                                                                                                      Score 78; DB 10; Length 25; Pred. No. 0.0038; Indels 1; Indels
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US-10-268-561-11

VS-10-268-561-11

Sequence 11, Application US/10268561

Publication No. US20030148333A1

GENERAL INFORMATION:

APPLICANT: Ortho-Clinical Diagnostics, Inc.

FIILE REFERENCE: CDS0286

FILE REFERENCE: CDS0286

CURRENT APPLICATION NUMBER: US/10/268,561

CURRENT APPLICATION NUMBER: US/10/268,561

CURRENT FILING DATE: 2002-10-10

PRIOR APPLICATION UNUBER: US 60/337453

PRIOR FILING DATE: 2001-11-05

NUMBER OF SEQ ID NOS: 19

SOFTWARE: PatentIn version 3.1

LENGTH: 15

Type: Det
RESULT 10
US-10-268-569-11
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Best Local S
Matches 13
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                                                                                                                                        Matches
                                                                                                                                                      Query Match
Best Local
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APPLICANT: Stewart, James L.
TITLE OF INVENTION: METHODS FOR THE SIMULTANEOUS DETECTION
TITLE OF INVENTION: OF HCV ANTIGENS AND HCV ANTIBODIES
FILE REFERENCE: 6821.US.P1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Abbott Laboratories APPLICANT: Shah, Dinesh O.
                                                                                                                                                                                                                                             TYPE: PRT
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                                                                    1 PRGSRPSWGPT 11
|||||||||
5 PRGSRPSWGPT 15
                                                                                                                                        11;
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Gutierrez, F
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Desai, Suresh
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Sequence 11, Application US/10268569
Publication No. US20030152965A1
GENERAL INFORMATION:
APPLICANT: Ortho-Clinical Diagnostics, Inc.
TITLE OF INVENTION: HCV Core Protein Sequen
FILE REFERENCE: CDS-0288

Sequences

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APPLICANT: Ortho-Clinical Diagnostics, Inc.
APPLICANT: Ortho-Clinical Diagnostics, Inc.
FILE OF INVENTION: HCV Core Protein Sequences
FILE REFERENCE: CDS-0288
CURRENT APPLICATION NUMBER: US/10/268,569
CURRENT FILLING DATE: 2002-10-10
PRIOR APPLICATION NUMBER: 60/347,303
PRIOR APPLICATION NUMBER: 60/347,303
PRIOR FILING DATE: 2001-11-11
NUMBER OF SEQ ID NOS: 19
SOFTWARE: Patentin version 3.1
SEQ ID NO 12
LENGTH: 15
TYPE: PRT
ORGANISM: Hepatitis C virus
US-10-268-569-12
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Sequence 12, Application US/10268569
Publication No. US20030152965A1
GENERAL INFORMATION:
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US-10-268-561-12
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Publication No. US20030148333A1
GENERAL INFORMATION:
APPLICANT: Ortho-Clinical Diagnostics, Inc.
TITLE OF INVENTION: HCV Anti-Core Monoclonal
FILE REFERENCE: CDS0286
FILE REFERENCE: CDS0286
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CURRENT FILING DATE: 2002-10-10
PRIOR APPLICATION NUMBER: 60/347,303
PRIOR FILING DATE: 2001-11-11
NUMBER OF SEQ ID NOS: 19
SOFTWARE: Patentin version 3.1
SEQ ID NO 11
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Matches
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SEQ ID NO 12
LENGTH: 15
                Query Match
Best Local 9
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Best Local
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CURRENT FILING DATE: 2002-10-10
PRIOR APPLICATION NUMBER: US 60/337453
PRIOR FILING DATE: 2001-11-05
NUMBER OF SEQ ID NOS: 19
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TYPB: PRT
ORGANISM: Hepatitis C virus
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52.8%; ilarity 91.7%; Conservative
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100.0%;
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91.7%;
Score 65; DB 14
Pred. No. 0.11;
0; Mismatches
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Pred. No.
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                                    DB 14; Length 15;
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Sequence 180, Application US/10350719

Publication No. US20030219726A1

GENERAL INFORMATION:
APPLICANT: Doorbar, John

ITILE OF INVENTION: IMPROVEMENTS IN OR RELATING TO TITLE OF INVENTION: VIRUSES

FILE REFERENCE: 18396/2162

CURRENT FILING DATE: 2003-01-24

PRIOR APPLICATION NUMBER: US/10/350,719

CURRENT FILING DATE: 2001-03-16

PRIOR FILING DATE: 2001-03-16

PRIOR APPLICATION NUMBER: GENOO18140.4

PRIOR FILING DATE: 2000-07-24

NUMBER OF SEQ ID NOS: 180

SOFTWARE: Patentin Ver. 2.1

SEQ ID NO 180

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PRIOR FILING DATE: 1997-11-19
PRIOR PELICATION NUMBER: PCT/EP94/0355
PRIOR PILING DATE: 1994-10-28
PRIOR APPLICATION NUMBER: EP 93402718.6
PRIOR FILING DATE: 1993-11-04
NUMBER OF SEQ ID NOS: 286
VOMBER OF SEQ ID NOS: 286
PRIOR PRIOR PLING DATE: 1993-11-04
NUMBER OF SEQ ID NOS: 286
PRIOR P
                                                                                                                                                                                                      PEATURE:

NAME/KEY: misc feature

LOCATION: (18)...(18)

OTHER INFORMATION: Xaa can be any naturally occurring amino
US-10-350-719-180
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US-10-651-165-9
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Publication No. US20040047877A1
                                                                                                           Query Match
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APPLICANT: DELEYS, Robert
APPLICANT: MAERTENS, Geert
TITLE OF INVENTION: IMMUNODOMINANT HUMAN T
TITLE OF INVENTION: VIRUS
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CURRENT APPLICATION NUMBER: US/10/651,165
CURRENT FILING DATE: 2003-09-02
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: PRT ORGANISM: Human papillomavirus
Local Similarity tes 8; Conserv
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nilarity 57.1%;
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                Score 54; DB Pred. No. 3.3; 1; Mismatches
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1.5;
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2: pir2:*
3: pir3:*
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144.287 Million cell updates/sec
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I lig	hypothetical prote	ribosomal protein	5	outer laver protei	inase			hypothetical prote			protein	μ.	01.	- rain	protamine CII - ch

#### ALIGNMENTS

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orf 3' of cycl - Rhodobacter sphaeroides (fragment)
C;Species: Rhodobacter sphaeroides
C;Date: 21-Sep-1993 #sequence_revision 18-Nov-1994 #text_change 18-Nov-1994
C;Accession: C40638
R;Rott, M.A.; Witthuhn, V.C.; Schilke, B.A.; Soranno, M.; Ali, A.; Donohue, T.J.
J. Bacteriol. 175, 358-366, 1993
A;Title: Genetic evidence for the role of isocytochrome c2 in photosynthetic gro
A;Reference number: A40638; MUID:93123153; PMID:8380401
A;Accession: C40638
A;Status: preliminary
A;Status: preliminary
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C40638
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                                                                                                                                                             A;Molecule type: nucleic acid
A;Residues: 1-29 <ROT>
A;Note: sequence extracted fr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A;Molecule type: mRNA
A;Residues: 1-23 <PAG-
A;Crose-references: EMBL:X52157; NID:g59010; PIDN:CAA36410.1; PID:g59011
C;Superfamily: coronavirus nucleocapsid protein
C;Keywords: nucleocapsid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                nucleocapsid protein - porcine transmissible gastroenteritis virus (fragment) (;Species: porcine transmissible gastroenteritis virus (;Date: 31-Dec-1990 #sequence_revision 31-Dec-1990 #text_change 20-Sep-1999 C;Accession: S10448 R;Page, K.W.; Britton, P.; Boursnell, M.E.G. submitted to the EMBL Data Library, March 1990 A;Reference number: S10447 A;Reference number: S10447 A;Reference number: S10447
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                                                                                                   Similarity
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QGFRPRHGATGDRHTLGLKRDRSV 28
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                                                                                                                                                      extracted from NCBI backbone (NCBIN:122345, NCBIP:123661)
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2; Mismatches
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RESULT B70566

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myrB protein - Micromonospora griseorubida (fragment)
()Species: Micromonospora griseorubida
()Date: 19-Mar-1997 #sequence_revision 18-Jul-1997 #text_change 05-Nov-1999
()Accession: $51593
R. Inouye, M.; Takada, Y.; Muto, N.; Beppu, T.; Horinouchi, S.
Mol. Gen. Genet. 245, 456-464, 1994
A;Title: Characterization and expression of a P-450-like mycinamicin biosynthesis gene
A;Reference number: $51593; MUID:95107242; PMID:7808395
A;Accession: $51593; MUID:95107242; PMID:7808395
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(;Species: Mycobacterium tuberculosis
(;Date: 17-Jul-1998 #sequence_revisoin 17-Jul-1998 #text_change 20-Jun-2000
(;Accession: B70566
R;Cole, S.T.; Brosch, R.; Parkhill, J.; Garnier, T.; Churcher, C.; Harris, D.; Gordon, Connor, R.; Davies, R.; Devikin, K.; Feltwell, T.; Gentles, S.; Hamlin, N.; Holroyd, Connor, R.; Davies, R.; Devikin, K.; Selseger, K.; Skelton, S.; Squares, S.
Rajandzeam, M.A.; Rogers, J.; Rutter, S.; Seeger, K.; Skelton, S.; Squares, S.
Nature 393, 537-544, 1998
A;Authors: Sqares, R.; Sulston, J.E.; Taylor, K.; Whitehead, S.; Barrell, B.G.
A;Title: Deciphering the biology of Mycobacterium tuberculosis from the complete genome A;Reference number: A70500; MUID: 98295987; PMID: 9634230
A;Accession: B70566
                       A;Cross-references: EMBL:D16098; NID:g286050; PIDN:BAA03671.1; PID:d1004186; PID:g829046
                                                     A; Molecule type: DNA
A; Residues: 1-28 < INO>
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A; Gene: merD
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C;Species: Yersinia enterocolitica
C;Date: 06-Jan-1995 #sequence_revision 26-Jul-1996 #text_change 08-Oct-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Ş
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C;Superfamily: Escherichia coli ribosomal protein L36
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A;Experimental source: strain H37Rv
C;Genetics:
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A;Molecule type: DNA
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A;Residues: 1-24 <OSB>
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Genetics:
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les 5; Conservation
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7; Conserve
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43.8%; Pred. No. 26
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hypothetical protein SC10H5.05 SC10H5.05 - Streptomyces coelicolor C;Species: Streptomyces coelicolor C;Date: 05-Nov-1999 #sequence_revision 05-Nov-1999 #text_change 05-Nov-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   R:Cole, S.T.; Biglmeier, K.; Parkhill, J.; James, K.D.; Thom R: Davies, R.M.; Devlin, K.; Duthoy, S.; Feltwell, T.; Frasem, M.A.; Rutherford, K.m. ature 409, 1007-1011, 2001

Nature 409, 1007-1011, 2001

A;Authors: Rutter, S.; Seeger, K.; Simon, S.; Simmonds, M.; A;Authors: Rutter, S.; Seeger, M.; Deprosy bacillus.

A;Reference number: A86909; MUID:21128732; PMID:11234002
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C;Superfamily: Escherichia coli ribosomal protein L36
                                                                    T34643
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       B
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Int. J. Protein Res. 1, 221-224, 1969
Int. J. Protein Res. 1, 221-224, 1969
A;Title: A new method for fractionation of protamines and the amino acid sequences of sa A;Reference number: A91774; MUID:72030546; PMID:4953052
A;Accession: A02670
                                                                                                                                                                                                                                                                                                                                                                                              A;Cross-references: GB:AL450380; NID:g13093616; PIDN:CAC30916.1; GSPDB:GN00147
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A; Residues: 1-37 < STO>
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C;Date: 12-Aug-1981 #sequence_revision 12-Aug-1981 #text_change 16-Feb-1997
C;Accession: A02670
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A; Residues: 1-33 < AND>
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N; Alternate nam
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K.; Duthoy, S.; Feltwell, T.; Fraser, A.; Hamlin, N.;
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Pred. No. 2.8e+02;
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Pred. No.
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A;Cross=references: GB:X59250; GB:S81088; NID:g44071; PIDN:CAA41942.1; PID:g44076
A;Note: sequence extracted from NCBI backbone (NCBIN:81088; NCBIP:81093)
R;Bolotin, A.; Wincker, P.; Mauger, S.; Jaillon, O.; Malarme, K.; Weissenbach, J.; Ehrli Genome Res. 11, 731-753, 2001
A;Title: The complete genome sequence of the lactic acid bacterium Lactococcus lactis sg A;Reference number: A86625; MUID:21235186; PMID:11337471
A;Accession: G86883
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J. Gen. Microbiol. 137, 2595-2600, 1991
A;Title: Nucleotide sequence of a Lactococcus lactis gene cluster encoding A;Reference number: A44812; MUID:92148378; PMID:1783905
A;Accession: D44812
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C;Date: 17-Apr-1993 #sequence_revision 17-Apr-1993 #text_change 03-Aug-2001
C;Accession: S17989; D44812; G86883
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A; Residues: 1-38 <STO>
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A; Residues: 1-38 < KO2>
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A;Note: the nucleotide sequence was
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FEBS Lett. 288, 114-118, 1991
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9; Conserv
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Pred. No. 3.4e+02;
2; Mismatches 8;
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Mismatches 2;
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RESULT

A;Accession:

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C;Species: Streptococ
C;Date: 22-Oct-2001 †
C;Accession: D97898
R;Hoskins, J.A.; Albo
R;Maier, R.M.; Neckermann, K.; Igloi, G.L.; Koessel, H. 
J. Mol. Biol. 251, 614-628, 1995
A;Tille: Complete sequence of the maize chloroplast genome:
A;Reference number: S58531; MUID:95395841; PMID:7666415
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                                                                                                                                                                      hypothetical protein 38 - maize chloroplast
C;Species: chloroplast Zea mays (maize)
C;Date: 29-Nov-1995 #sequence_revision 19-Jan-1996 #text_
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D97898
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A;Reference number: A97872; MUID:21429245; PMID:11544234
A;Accession: D97898
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A; Residues: 1-38 < KUR>
A; Cross-references: GB
C; Genetics:
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C;Species: Streptococcus pneumoniae
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A; Residues: 1-38 < KUR>
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R.; LeBlanc, D.J.; Lee, L.N.; Lefkowitz, E.J.; Lu, J.; Matsushima, P.; McAhren,
P.; Sun, P. M.; Winkler, M.E.
Bacteriol. 183, 5709-5717, 2001
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Pred. No. 5.2e+02;
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A;Molecule type: DNA
A;Residues: 1-38 <MAI>
A;Residues: 1-38 <MAI>
A;Cross-references: EMBL:X86563; NID:g902200; PIDN:CAA60335.1; PID:g902270
A;Note: the nucleotide sequence was submitted to the EMBL Data Library, April 1995
C;Genetics:
                                                                                                                                              A,Molecule type: protein
A,Residues: 2-33 <MCX>
C;Superfamily: protamine Y2
C;Keywords: DNA binding; nucleus
F;2-33/Product: protamine TP21 #status experimental <MAT>
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A;Residues: 1-33 <AIK>
A;Residues: 1-33 <AIK>
A;Cross-references: EMBL:X01599
A;Cross-references: EMBL:X01599
R;McKay, D.J.; Renaux, B.S.; Dixon, G.H.
Eur. J. Blochem. 158, 361-366, 1986
Eur. J. Blochem. 158, 361-366, 1986
A;Title: Rainbow trout proteamines. Amino acid sequences of six distinct proteins from A;Reference number: A91170; MUID:86274711; PMID:3755398
A;Accession: C24970
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Note: Incomplete the protamine and protamine TP21 - rainbow trout)

C;Species: Oncorthynchus mykiss (rainbow trout)

C;Date: 30-Jun-1992 #sequence revision 30-Jun-1992 #text_change 01-Aug-1997

C;Accession: E21211; C24970

C;Accession: E21211; C24970

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A;Title: Sequence homologies in the protamine gene family of rainbow trout.
A;Reference number: A21211; MUID:83272939; PMID:6308564
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A;Molecule type: DNA
A;Residues: 1-33 <AIK>
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3; Mismatches
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Pred. No. 6.2e+02;
1; Mismatches 9
     Score 31;
Pred. No.
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J. 5.2e+02;
J. 3.2 9;
           DB 2; Length 33; 6.2e+02;
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G;Species: Pseudomonas sp.

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G;Date: 31-Dec-1993 #sequence_revision 31-Dec-1993 #text_change 21-Mar-1996
G;Accession: A42592
R;Blanche, F:, Thibaut, D.; Famechon, A.; Debussche, L.; Cameron, B.; Crouzet, J.
J. Bacteriol. 174, 1036-1042, 1992
A;Title: Precorrin-6x reductase from Pseudomonas denitrificans: purification and charact A;Reference number: A42592; MUID:92121090; PMID:1732193
A;Accession: A42592
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MEDLINE=22242508; Publ
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AAM78670.1; -...
BAC63160.1; -...
AAL96899.1; -...
CAD45726.1; -...
AAM98989.1; -...
AAM58607.1; -...
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k C., Buchrieser
M., Couve E., La
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PubMed=12200547;
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          Complete proteome.
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the L36P family of ribosomal proteins.
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ui L., Poyart
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A., Ferretti
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A Cole S.T., Brosch R., Parkhill J., Garnier T., Churcher C., Harrii A. Cole S.T., Brosch R., Parkhill J., Garnier T., Churcher C., Harrii A. Gordon S.V., Elglmeier K., Gas S., Barry C.E. III, Tekaia F., A. Badcock K., Basham D., Brown D., Chillingworth T., Connor R., Davies R., Devlin K., Feltwell T., Gentles S., Hamlin N., Holroyd A. Hornsby T., Jagels K., Krogh A., McLean J., Moule S., Murphy L., A. Coliver S., Osborne J., Quail M.A., Rajandream M.A., Rogers J., A. Rutter S., Seeger K., Skelton S., Squares S., Squares R., Rutter S., Seeger K., Skelton S., Squares S., Squares R., Sulston J.E., Taylor K., Whitehead S., Barrell B.G.; Sulston J.E., Taylor K., Whitehead S., Barrell B.G.; Squares S., Squares S
                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE FROM N.A.
SPECIES=M.Bovis; STRAIN=BCG;
SPECIES=M.Bovis; STRAIN=BCG;
MEDLINE=96200849; PubMed=8621083;
Dubnau E., Soares S., Huang T.J., Ja
Hubnau E., Soares S., Huang T.J., Ja
Mycobacterium smegmatis.";
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SPECIES=M.tuberculosis; STRAIN=CDL 13.1. ,

MEDLINE=22206494; PubMede12218036;

MEDLINE=22206494; PubMede12218036;

MEDLINE=22206494; PubMede12218036;

Pleischmann R.D., Alland D., Eisen J.A., Carpenter L., Hickey E.,

Peterson J., DeBoy R., Dodson R., Gwinn M., Haft D., Hickey E.,

Kolonay J.F., Nelson W.C., Umayam L.A., Ermolaeva M., Salzberg S.

Kolonay J.F., Nelson W.C., Umayam L.A., Ermolaeva M., Salzberg S.

Delcher A., Utterback T., Weidman J., Khouri H., Gill J., Mikula

Bishai W., Jacobs W.R. Jr., Venter J.C., Fraser C.M.;

Bishai W., Jacobs W.R. Jr., Venter J.C., Fraser C.M.;
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01-NOV-1995 (Rel. 32, Created)
01-NOV-1995 (Rel. 32, Last sequence update)
10-CCT-2003 (Rel. 42, Last annotation update)
50S ribosomal protein L36 (Ribosomal protein
RPMJ OR RV3461C OR MT3567.1 OR MTCY13E12.14C
                                                                    SPECIES—N. Dovis; STRAIN=AF2122/97;

SPECIES—N. Dovis; STRAIN=AF2122/97;

MEDLINE=22709107; PubMed=12788972;

MEDLINE=22709107; PubMed=12788972;

Marier T., Eiglmeier K., Camus J.-C., Medina N., Mansoor H.,

Procr M., Duthoy S., Grondin S., Lacroix C., Monsempe C., Simon S.

Harris B., Atkin R., Doggett J., Mayes R., Keating L., Wheeler P.:

Parkhill J., Barrell B.G., Cole S.T., Gordon S.V., Hewinson R.G.,

"The complete genome sequence of Mycobacterium bovis.";

Proc. Natl. Acad. Sci. U.S.A. 100:7877-7882(2003).

Proc. Natl. Acad. Sci. U.S.A. 100:7877-7882(2003).
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MEDLINE=98295987; PubMed
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Corynebacterineae, Mycobacteriaceae, Mycobacterium.
NCBI_TaxID=1773, 1765;
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                           This
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SWISS-PROT entry is copyright. It is produced sen the Swiss Institute of Bioinformatics and
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28-FEB-2003
                                                                                                                                                                                                                               This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Salanoubat M., Genin S., Artiguenave F., Gouzy J., Mangenot S., Arlat M., Billault A., Brottier P., Camus J.C., Cattolico L., Chandler M., Choisne N., Claudel-Renard C., Cunnac S., Demange N., Gaspin C., Lavie M., Moisan A., Robert C., Saurin W., Schiex T., Siguier P., Thebault P., Whalen M., Wincker P., Levy M., Weissenbach J., Boucher C.A., "Genome sequence of the plant pathogen Ralstonia solanacearum.", "Genome seq
                                                                                             EMBL; AL646073; CAD16706.1; -. HAMAP; MF_00251; -; 1. InterPro; IPR000473; Ribosomal_L36.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             28-FEB-2003 (Rel. 41, Created)
28-FEB-2003 (Rel. 41, Last sequence update)
28-FEB-2003 (Rel. 41, Last annotation update)
28-FEB-2003 (Rel. 41, Last annotation update)
50-FEB-2003 (Rel. 41, Last annotation update)
50-FEB-2003 (Rel. 41, Last annotation update)
50-FEB-2003 (Rel. 41, Last sequence update)
60-FEB-2003 (Rel. 41, Last 
                           Pfam; PF00444; Ribosomal L36; ProDom; PD002101; Ribosomal L
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDLINE=21681879; PubMed=11823852;
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ProDom; PD002101; Ribosomal L36; 1.
TIGRPAMS; TIGR01022; rpmJ_bact; 1.
PROSITE; PS00828; RIBOSOMAL L36; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Burkholderiaceae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Ribosomal protein; Complete proteome.
SEQUENCE 37 AA; 4343 MW; C92A4845315F930F CRC64;
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HSSP; P80256; 1DFE.
TIGR; MT3567.1; -.
Tuberculist; Rv3461c; -.
HAMAP; MF_00251; -; 1.
InterPro; IRR000473; Ribosomal_L36.
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EMBL; AE007160; AAK47907.1; -.
EMBL; U15140; AAB17596.1; -.
EMBL; BX248346; CAD95677.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            entities requires a license agreement (See http://www.isb-sib.ch/announce) or send an email to license@sisb-sib.ch).
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5; Conserv
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TIGR01022;
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Pred. No. 1.1e+02;
2; Mismatches 1
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Best Local S
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Q9X7A2;
30-MAY-2000
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Ando T, Watanabe S.;

"A new method for fractionation of protamines and the amino acid sequences of salmine and three components of iridine.";

Int. J. Protein Res. 1:221-224 (1969).

-!- FUNCTION: Protamines substitute for histones in the chromatin of sperm during the haploid phase of spermatogenesis. They compact sperm DNA into a highly condensed, stable and inactive complex.

-!- SUBCELTULIAR LOCATION: Nuclear.

-!- TISSUE SPECIFICITY: Testis.
MEDLINE-21128732; PubMed=11234002;
Cole S.T., Eiglmeier K., Parkhill J., James K.D., Thomson N.R.
Wheeler P.R., Honore N., Garnier T., Churcher C., Harris D.,
Mungall K., Basham D., Brown D., Chillingworth T., Connor R.,
Davies R.M., Devlin K., Duthoy S., Feltwell T., Fraser A., Ha
                                                                                                                                                                                                                                                                                                                       30-MAY-2000 (Rel. 39, Created)
30-MAY-2000 (Rel. 39, Last sequence update)
28-FEB-2003 (Rel. 41, Last annotation update)
50S ribosomal protein L36.
RPMJ OR ML1961 OR MLCB1222.31C.
Mycobacterium leprae.
Bacteria; Actinobacteria; Actinobacteridae; Actinomycetales;
Corynabacterineae; Mycobacteriaceae; Mycobacterium.
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P02328;
                                                                                                                                                                                                                                SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                             NCBI_TaxID=1769;
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; E
Actinopterygii; Neopterygii; Teleostei; Euteleostei;
Protacanthopterygii; Salmoniformes; Salmonidae; Oncor
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21-UTL-1986 (Rel. 01, Last sequence up
30-WAY-2000 (Rel. 39, Last annotation
Protamine IA (Iridine IA).
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condensation; Nuclear protein.
33 AA; 4406 MW; F82COA9C80A1FFF2 CRC64;
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Pred. No.
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); Mismatches
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                                                                                                                  Thomson N.R.,
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           Hamlin
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C P277
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                                                                                                         MEDLINE=92148378; PubMed=1783905;
Koivula T., Hemilae H.;
"Nucleotide sequence of a Lactococcus lactis gene cluster encoding
ademylate kinase, initiation factor I and ribosomal proteins.";
J. Gen. Microbiol. 137:2595-2600(1991).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               HAMAP; MF_00251; -; 1.
InterPro; IPR000473; Ribosomal L36; I.
Pfam; PF00444; Ribosomal L36; I.
TIGRPAMs; TIGR01022; rpmJ bact; 1.
PROSITE; PS00828; RIBOSOMĀL_L36; 1.
RIBOSOMĀL PSOUCEAIN; COMPLETE PROTECME.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Bolotin A., Wincker P., Mauger S., Jaillon O., Malarme K., Weissenbach J., Ehrlich S.D., Sorokin A.; Weissenbach genome sence of the lactic acid bacterium Lactococcus "The complete genome encode of the lactic acid bacterium Lactococcus lactis ssp. lactis II1403."; Genome Res. 11:731-753(2001).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Holroyd S., Hornsby T., Jagels K., Lacroix C., Maclean J., Moule S., Murphy L., Oliver K., Quail M.A., Rajandream M.A., Rutherford K.M., Rutter S., Seeger K., Simon S., Simonds M., Skelton J., Squares R., Squares S., Stevens K., Taylor K., Whitehead S., Woodward J.R., Macrill B.G.;
           SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SPECIES=L.1.lactis; STRAIN=IL1403; MEDLINE=21235186; PubMed=11337471;
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EMBL; AL583923;
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HSSP; P80256; 1DFE.
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                                                                                                                                                                                                                                                                                                                                                                                       cremoris; STRAIN=MG1614;
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ilarity 71.4%;
Conservative
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Pred. No. 1.5e
1; Mismatches
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Best Local
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PIR; D97898; D97898.

PIR; S17989; S17989.

HSSP; P80256; 1DFE.

TIGR; SP0233; -..

InterPro; IPR000473; Ribosomal_L36.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE FROM N.A.

SPECIES-S.pneumoniae; STRAIN=ATCC BAA-255 / R6;

MEDLINE-214.9245;

MEDLINE-214.9245;

PubMed=11544234;

MEDLINE-214.9245;

PubMed F. Jr., Arnold J., Blaszczak L.C., Burgett S. DeHoff B.S., Estrem S.T., Fritz L., Fu D.-J., Fuller W., Geringer DeHoff B.S., Strem S.T., Khoja H., Kraft A.R., Lagace R.E., Gilmour R., Glass J.S., Khoja H., Kraft A.R., Lagace R.E., LeBlanc D.J., Lee L.N., Lefkowitz E.J., Lu J., Matsushima P., McBahern S.M., McHenney M., McLeaster K., Mundy C.W., Nicas T.I., Nochen, McHenney M., McLeaster K., Mundy C.W., Nicas T.I., Norris F.H., O'Gara M., Peery R.B., Robertson G.T., Rockey P., Sun P.-M., Winkler M.E., Yang Y., Young-Bellido M., Zhao G., Zook C.A., Baltz R.H., Jaskunas S.R., Rosteck P.R. Jr., Skatrud P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SPECIES=S.pneumoniae; STRAIN=ATCC BAA-334 / TIGR4;
MEDLINB=21357209; PubMed=11433916;
Tettelin H., Nelson K.E., Paulsen I.T., Eisen J.A., Read T.D.,
Peterson S., Heidelberg J., DeBoy R.T., Haft D.H., Dodson R.J.,
Durkin A.S., Gwinn M., Kolonay J.F., Nelson W.C., Peterson J.D.,
Umayam L.A., White O., Salzberg S.L., Iewis M.R., Radune D.,
Holtzapple E., Khouri H., Wolf A.M., Utterback T.R., Hansen C.L.,
McDonald L.A., Feldblyum T.V., Argiuoli S., Dickinson T., Hickey E.K.,
McDonald L.A., Feldblyum T.V., Argiuoli S., Dickinson T., Hickey E.K.,
Dougherty B.A., Morrison D.A., Hollingshead S.K., Fraser C.M.;
"Complete genome sequence of a virulent isolate of Streptococcus
                                                                                                                                                            SEQUENCE
                                                                                                                                                                                                 TIGRFAMs; TIGR01022; rpmJ bact; 1. PROSITE; PS00828; RIBOSOMAL_L36; 1.
                                                                                                                                                                                                                  TIGRFAMs;
                                                                                                                                                                                                                                      Pfam; PF00444; Ribosomal_L36; 1.
ProDom; PD002101; Ribosomal_L36; 1.
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EMBL; AE008402; AAK99016.1; -.
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MEDLINE=91348245; PubMed=1908794;
Moivula T., Palva I., Hemilae H.;
"Nucleotide sequence of the secY gene from Lactococcus lactis and identification of conserved regions by comparison of four SecY proteins.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Glass J.I.;
"Genome of the bacterium Streptococcus pneumoniae strain R6.";
J. Bacteriol. 183:5709-5717(2001).
-!- SIMILARITY: Belongs to the L36P family of ribosomal proteins.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Science 293:498-506(2001).
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28
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                                                                                               Similarity
PANPKHKQR 36
                                                                                                                                                        protein;
                                       PTDPRHRSR
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                                                                                                                                                          Complete proteome. 4421 MW; 892FEFE9A5939D88 CRC64;
                                    18
                                                                                             26.0%;
44.4%;
                                                                                               Score 32;
Pred. No.
                                                                                                                       DB 1;
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RESULT 8
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P08146;
01-AUG-1988 (Rel. 0
01-AUG-1988 (Rel. 0
15-DBC-1998 (Rel. 3
                                                                                                                                                                                                                                                                      PRT8_ONCMY
P12817;
01-OCT-1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDLINE-83272939; PubMed-6308564;
Aiken J.M., McKenzie D., Zhao H.-Z., States J.C., Dixon G.H.;
Aiken J.M., McKenzie D., Zhao H.-Z., States J.C., Dixon G.H.;
Nucleic Acids Res. 11:4907-4922(1983).
-!- FUNCTION: Protamines substitute for histones in the chromatin of sperm during the haploid phase of spermatogenesis. They compact sperm DNA into a highly condensed, stable and inactive complex.
-!- SUBCELLULAR LOCATION: Nuclear.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial
                                                                                                                                              Oncorhynchus mykiss (Rainbow trout) (Salmo gairdneri).
Eukaryota; Metazoa; Chordata; Craniata; Verrebrata; Euteleostomi;
Actinopterygii; Neopterygii; Teleostei; Euteleostei;
Protacanthopterygii; Salmoniformes; Salmonidae; Oncorhynchus.
NCBI_TaxID=8022;
                                                                                                                                                                                                                                                                                                                   ONCMY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 entities requires a license agreement (See http:\overline{//www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Actinopterygii; Neopterygii; Teleostei; Euteleostei;
"Rainbow trout protamines. Amino acid sequences of six d proteins from a single testis."; Eur. J. Biochem. 158:361-366(1986).
-!- FUNCTION: Protamines substitute for histones in the sperm during the haploid phase of spermatogenesis. I sperm DNA into a highly condensed, stable and inacti
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE FROM N.A.
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01-APR-1993
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                                                                                          McKay
                                                                                                      MEDLINE=86274711; PubMed=3755398;
                                                                                       D.J., Renaux B.S., Dixon G.H.;
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(Rel. 25, Last annotation updat
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R HSSP; P80256; 1DFE.

R HSSP; H10798.1; -.

R HAMAP; MF 00251; -; 1.

R InterPro; IPR000473; Ribosomal L36.

R Pfam; PF00444; Ribosomal L36; I.

R ProDom; PD002101; Ribosomal L36; 1.

R PTORFAMS; TIGR01022; rpmJ bact; 1.

R PROSITE; PS00828; RIBOSOMAL L36; 1.

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01-NOV-1995
01-NOV-1995
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                                                                                                                                                                                                                                                                                              This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/
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01-NOV-1995 (Rel. 32, Last sequence up
28-FEB-2003 (Rel. 41, Last annotation
50s ribosomal protein L36.
EPMJ OR RPL36 OR HI0798.1.
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-!- TISSUE SPECIFICITY: Testis.
PIR; E21211; E21211.
                                                                                                                                                                                                                                                                               entities requires a license agreement (Some send an email to license@isb-sib.ch).
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MEDLINE=95350630; PubMed=7542800;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Chromosomal protein; Nucleosome core; Spermatogenesis; DNA-binding; Testis; DNA condensation; Nuclear protein. SEQUENCE 32 AA; 4224 MW; B07A9C9D90B796FF CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                             Science 269:496-512(1995).
-!- SIMILARITY: Belongs to the L36P family of ribosomal proteins.
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STRAIN=Rd / KW20 / ATCC 51907;
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Bacteria; Proteobacteria; Gam
Pasteurellaceae; Haemophilus.
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Gomes C.M., Faria A., Carita J., Mendes J.C., Huber H., Stetter K.O., Teixeira M.; "Di-cluster, seven iron ferredoxins from hyper Sulfolbales.";
J. Biol. Inorg. Chem. 3:499-507(1998).
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P81542;
15-JUL-1999
15-JUL-1999
10-OCT-2003
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HSSP; P80251; -; 1.
HAMAP; MF 00251; -; 1.
InterPro; IPR000473; Ribosomal L36; 1.
Pfam; PF00444; Ribosomal L36; 1.
Procom; PD002101; Ribosomal L36; 1.
Pfam; Pf00447; PfmJ_bact; 1.
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P57942;
16-OCT-2001 |
16-OCT-2001
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ProDom; PD002101; Ribosomal 136; 1.
TIGREAMS; TIGR01022; rpmJ bact; 1.
PROSITE; PS00828; RIBOSOMAL 136; 1.
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16-OCT-2001 (Rel. 40, Last sequence update)
28-FBB-2003 (Rel. 41, Last annotation update)
50S ribosomal protein L36.
REMJ OR RFL36 OR RMI394.
                                                                                                                                                SEQUENCE.
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-!- SIMILARITY: Belongs to the L36P family of ribosomal proteins.
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Pasteurellaceae; Pasteurella.
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(Rel. 42, Last annotation update)
ning ferredoxin (Seven-iron ferredoxin)
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21-JUL-1986 (Rel. 0
01-NOV-1988 (Rel. 0
28-FEB-2003 (Rel. 4
                                                                                   Bombesin
PEPTIDE
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"Neuromedin B: a novel bombesin-like peptide identified in porcine spinal cord.";
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Minamino N., Sudoh T., Kangawa K., M
"Neuromedin B-32 and B-30: two 'big'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Eukaryota; Metazoa;
Mammalia; Eutheria;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 METAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       -
                                                                                                                                                                                                                                                                                                           ÷
                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE OF 23-32.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               -!- FUNCTION:
                                                                                                                                                                          nterPro;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           porcine brain and spinal cord.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            METAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DOMAIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Electron transport; Iron-sulfur; Repeat; 4Fe-4S; 3Fe-4S; Methylation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       issp;
                                                                                                                                                                                                                                                             chem. Biophys. Res. Commun. 114:541-548(1983).
FUNCTION: Stimulates smooth muscle contraction in a manner similar
to that of bombesin.
SUBCELLULAR LOCATION: Secreted.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FUNCTION: Ferredoxins are iron-sulfur proteins that transfer electrons in a wide variety of metabolic reactions. COFACTOR: Binds one 3Fe-4S and one 4Fe-4S cluster and one zinc
                                                                                                                                                                                                                                        SIMILARITY: Belongs to the bombesin/neuromedin B/ranatensin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SIMILARITY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               atom (By similarity)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PIG
                                                                                                                                                                                                  B60301; BSPGNB.
                                                                                                                                                   PF02044; Bombesin;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    P55907; 1XER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2 RGSRPSWG------PTDPRHRSRNVG 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Similarity 9; Conser
                                                                                          family; Amidation.
23 32
                                                                                                                                  PS00257;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RTSRPVTGDHAGHKVYAPADPPVKEKALG 35
                                                                                                                                                                          IPR000874; Bombesin.
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                                                                                                                                  BOMBESIN; 1.
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41,
                                          3654 MW;
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24.4%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 sequence update)
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Score 30;
                                          7B3FC292C953DE80 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         256DB78618BC9ED5 CRC64;
                                                                  AMIDATION
                                                                                       NEUROMEDIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Pred. No. 3.5e+02
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         N-TERMINAL EXTENSION.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matsuo H.;
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DB 1;
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Length 32;
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                                                                                                                                                                                                                                                          HSSF; RB0256; IDFE.
HSSF; RB0256; IDFE.
THERPOOLOGY RABOSOMAL L36.
Ffam; PF00444; Ribosomal L36; I.
ProDom; PD002101; Ribosomal L36; I.
TIGREAMS; TIGR01022; rpmJ bact; 1.
PROSITE; PS00828; RIBOSOMĀL L36; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 This SWISS-PROT entry is copyright. It is produced through a collaborate between the Swiss Institute of Bioinformatics and the EMBL outstatist the European Bioinformatics Institute. There are no restrictions on use by non-profit institutions as long as its content is in no modified and this statement is not removed. Usage by and for commer entities requires a license agreement (See http://www.isb-sib.ch/annou
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDLINE=22598143; PubMed=12712204;

Ren S.-X., Fu G., Jiang X.-G., Zeng R., Miao Y.-G., Xu H.,

Zhang Y.-X., Xiong H., Lu G., Lu L.-F., Jiang H.-Q., Jia J., Tu Y.-F.

Jiang J.-X., Gu W.-Y., Zhang Y.-Q., Cai Z., Sheng H.-H., Yin H.-F.,

Zhang Y., Zhu G.-F., Wan M., Huang H.-L., Qian Z., Wang S.-Y., Ma W.,

Yao Z.-J., Shen Y., Qiang B.-Q., Xia Q.-C., Guo X.-K., Danchin A.,

Saint Girons I., Somerville R.L., Wen Y.-M., Shi M.-H., Chen Z.,

Xu J.-G., Zhao G.-P.;
                                                                                                                                                                                  Ribosomal protein; Complete proteome.

CONFLICT 24 24 G -> R (IN REF. 2
SEQUENCE 37 AA; 4182 MW; 5FC638666BF65D4E
                                                                                                                                                                                                                                                                                                                                                                                                                          EMBL; AF115283; AAD40606.1; -.
EMBL; AE011262; -; NOT_ANNOTATED_CDS.
EMBL; AE011263; -; NOT_ANNOTATED_CDS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE FROM N.A.
STRAIN-Lai / Serogroup Icterohaemo:
MEDLINE-2008835; PubMed=10620683;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LEPIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       or send
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Nature 422:888-893(2003).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       "Unique physiological and pathogenic features of Leptospira interrogans revealed by whole-genome sequencing.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE FROM N.A.
STRAIN=56601 / Serogroup Icterchaemorrhagiae / Serovar lai;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Zuerner R.L., Hartske
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Bacteria; Spirochaetes; Spirochaetales; Leptospiraceae; Leptospira.
NCBI_TaxID=173;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RPMJ OR LA0761.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FEMS Microbiol. Lett. 182:303-308(2000).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Leptospira interrogans.
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30-MAY-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           10-MAY-2000
10-OCT-2003
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                                                                                            4;
                                                                                                                    Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  s requires a license agreement (some email to license@isb-sib.ch).
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                                           TDPRHRSR 18
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(Rel. 39,
(Rel. 42,
                                                                                            Conservative
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                                                                                                               24.4%;
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                                                                                       Score 30; DB
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3; Mismatches
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4e+02;
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p57570;
p575701 (R
16-OCT-2001 (R
16-OCT-2003 (R
28-FEB-2003 (R
  STRAINETOKYO 1998;
MEDLINE=20445173; PubMed=10993077;
Shiqenobu S., Watanabe H., Hattori M., Sakaki Y.,
"Genome sequence of the endocellular bacterial sy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    EMBL; AP005369; BAC07655.1; -.

HAMAP; MF 00251; -; 1.

InterPro; IPR000473; Ribosomal L36.

Pf00444; Ribosomal L36; 1.

TIGRPANS; TIGR01022; rpmJ bact; 1.

PROSITE; PS00828; RIBOSOMAL L36; 1.

RIBOSOMAL D36; 1.
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Nakamura Y., Kaneko T., Sato S., Ikeuchi M., Katoh H., Sasamot
Watanabe A., Iriguchi M., Kawashima K., Kimura T., Kishida Y.,
Kiyokawa C., Kohara M., Matsumoto M., Matsuno A., Nakazaki N.,
Shimpo S., Sugimoto M., Takeuchi C., Yamada M., Tabata S.;
"Complete genome structure of the thermophilic cyanobacterium
Thermosynechococcus elongatus BP-1.";
DNA Res. 9:123-130(2002)
                                                                                                                                                                                      Enterobacteriaceae;
                                                                                                                                                                                                                                                           Buchnera aphidicola (subsp. Acyrthosiphon pisum) (Acyrthosiphon pisum
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SYNEL
                                                                                                              SEQUENCE FROM
                                                                                                                                                                                                               Bacteria; Proteobacteria;
                                                                                                                                                                                                                                       symbiotic bacterium)
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10-OCT-2003 (Rel. 42, Last sequence update)
10-OCT-2003 (Rel. 42, Last annotation update)
10-OCT-2003 (Rel. 42, Last annotation update)
508 ribosomal protein L36.
RPMJ OR RPLJ6 OR TSR0102.
Synechococcus elongatus (Thermosynechococcus elongatus Bacteria; Cyanobacteria; Chroococcales; Synechococcus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                -!- SIMILARITY: Belongs to the L36P family of ribosomal proteins.
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al protein L36.
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Matches 4; Conserv
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RISSP; P80256; 1DFE.

HAMAP; MF 00251; -; 1.

InterPro; IPR00473; Ribosomal L36.

Pfam; PF00444; Ribosomal L36; 1.

PFODOM; PD002101; Ribosomal L36; 1.

PFORPAMS; TIGR01022; rpmJ_bact; 1.

PROSITE; PS00828; RIBOSOMAL L36; 1.

RIBOSOMAl protein; Complete Proteome.

RIBOSOMAl protein; Complete Proteome.

SEQUENCE 38 AA; 4365 MW; 29F271414AA25621 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                     This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Buchnera sp. APS.";
Nature 407:81-86(2000).
Nature 407:817: Belongs to the L36P family of ribosomal proteins.
                                                                             30 DPKHKQR 36
                                                                                                12 DPRHRSR 18
                                                                                                                                                4; Conservative
                                                                                                                                                                    24.4%;
                                                                                                                                                  Score 30; DB 1; Length 38; Pred. No. 4.1e+02; 2; Mismatches 1; Indels
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Gapop 10.0 , Gapext 0.5
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Copyright (c) 1993 - 2004 Compugen Ltd.
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## ALIGNMENTS

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Created) Last sequence update) Last annotation update) (Fragment).	PRT; 33 AA.	RHSRNV 20	Score 40.5; DB 2; Length 39; Pred. No. 77; 3; Mismatches 5; Indels 7; Gaps 2;	9A35BCE0F0F5D258 CRC64;		transposon structures in soil bacteria."; col. 0:0-0(1999).					'96 SE13		Last annotation update)	Created)	PRT; 39 AA.	

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Inoue-Murayama M., Takenaki
"Origin and divergence of i
receptor genes.";
Primates 39:217-224 (1998)
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Eukaryota; Metazoa;
Mammalia; Eutheria;
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Coronavirus PRCV nucleoprotein, 5' region (Fragme
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Submitted (APR-2001) to the
EMBL; BC006692; AAH06692.1;
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Mus musculus (Mouse).
Eukaryota; Metazoa; Cl
Mammalia; Eutheria; R
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SEQUENCE FROM N.A.
STRAIN=PRCV 86/137004(British Isolate);
Page K.W., Britton P., Boursnell M.E.G.;
"Sequence analysis of the leader RNA of transmissible gastroenteritis virus and property of the control of the contr
                                                                                                                                                                                                                                                            Porcine respiratory coronavirus. Viruses; ssRNA positive-strand v Coronaviridae; Coronavirus.
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  Primates; Strepsirhini; Loridae; Nycticebus.
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Virus Genes 0:0-0(1990).
EMBL; XS2868; CAA36895.1; -.
PIR; S10448; S10448.
GO; GO:0019013; C:viral nucleocapsid; I1
Nucleocapsid.
NON TER 23
SEQUENCE 23 AA; 2565 MW; D761FC2797
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01-UN-2003 (TrEMBLrel. 24, Last ann
Coronavirus TGEV nucleoprotein, 5'r
Transmissible gastroenteritts virus
SEQUENCE FROM N.A.
STRAIN-RB50 / ATCC BAA-588;
MEDLINE=22827954; PubMed=12910271;
Parkhill J. Sebaihia M., Preston A.,
Harris D.E., Holden M.T.G., Churcher
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Q7WRA1
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Bacteria; Proteobacteria; E
Alcaligenaceae; Bordetella.
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STRAIN=TGEV FS772/70 (Virulent British I PAGE K.W., Britton P., Boursnell M.E.G. PAGE K.W., Britton P., Boursnell M.E.G. "Sequence analysis of the leader RNA of
                                                                                                                                                                                                                                                                                              RPMJ OR BB0053
                                                                                                                                                                                                                                                                                                                          50S ribosomal protein 136.
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EMBL; X52157; CAA36410.1; -.
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nilarity 38.9%;
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         Murphy L.D.,
C.M., Bentley
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Best Local S
Matches 6
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A Harris D.E., Holden M.T.G., Churcher C.M., Bentley S.D., Mungall K.L.,
A Harris D.E., Holden M.T.G., Churcher C.M., Bentley S.D., Mungall K.L.,
A Cerdeno-Tarraga A.M., Temple L., James K., Harris B., Quail M.A.,
Chillingworth T., Collins M., Cronin A., Davis P., Doggett J.,
A Achtman M., Atkin R., Baker S., Basham D., Bason N., Cherevach I.,
Chillingworth T., Collins M., Cronin A., Davis P., Doggett J.,
A Reltwell T., Goble A., Hamlin N., Hauser H., Holroyd S., Jagels K.,
A Feltwell T., Goble A., Hamlin N., Fauser H., Holroyd S., Jagels K.,
A Feltwell T., Goble B., Norberczak H., O'Neil S., Ormond D., Price C.,
A Feltwell T., Goble S., Norberczak H., O'Neil S., Crmond D., Price C.,
A Feltwell T., Sunders M., Saunders M., Saunders S., Stevens K.,
A Kabbinowitsch E., Rutter S., Sanders M., Squares R.,
A Sharp S., Simmonds M., Skelton J., Squares R.,
Comparative analysis of the genome sequences of Bordetella pertussis,
Unwin L., Whitehead S., Barrell B.G., Maskell D.J.;
Comparative analysis of the genome sequences of Bordetella pertussis,
Borderella parapertussis and Bordetella bronchiseptica.";
BMBL; BX640423; CAE39794.1;
M. Bibcornal World CAESTON CAESTO
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Achtman M., Atkin R., Baker S., Basham D., Bason N., Chorevach I.,
Achtlman M., Atkin R., Baker S., Basham D., Bason N., Chorevach I.,
Achtlman M., Atkin R., Baker S., Basham D., Bason N., Chorevach I.,
A Chillingworth T., Collins M., Cronin A., Davis P., Doggett J.,
A Leather S., Moule S., Norberczak H., O'Neil S., Ormond D., Jagels K.,
A Leather S., Moule S., Norberczak H., O'Neil S., Ormond D., Price C.,
A Rabbinowitsch E., Rutter S., Sanders M., Saunders D., Seeger K.,
A Sharp S., Simmonds M., Skelton J., Squares R., Squares S., Stevens K.,
Unwin L., Whitehead S., Barrell B.G., Maskell D.J.;
"Comparative analysis of the genome sequences of Bordetella pertussis,
Bordetella parapertussis and Bordetella bronchiseptica.";
Nat. Genet. 35:32-40(2003).
LEMBL, BA640437; CAB30555.1;
EMBL, BA640437; CAB30555.1;
CM Ribosomal protein; Complete proteome.
SBQUENCE 37 AA; 4332 MW; 53B3EBBE5487DADD CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
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01-OCT-2003 (TrEMBLrel. 25, C
01-OCT-2003 (TrEMBLrel. 25, L
01-OCT-2003 (TrEMBLrel. 25, I
50S ribosomal protein 136.
               Q7VTA9
Q7VTA9;
01-OCT-2003
01-OCT-2003
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STRAIN=12822 / ATCC BAA-587;
STRAIN=22827954; PubMed=12910271;
Preston /
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Bacteria; Proteobacteria; E
Alcaligenaceae; Bordetella.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               protein; Complete proteome.
37 AA; 4332 MW; 53B3EBBE5487DADD CRC64;
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nilarity 75.0%;
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Pred. No. 1.7e+02;
1; Mismatches 1;
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       sequence update)
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RA Parkhill J., Sebalhia M., Preston A., Murphy L.D., Thomson N.,

RA Harris D.E., Holden M.T.G., Churcher C.M., Bentley S.D., Mungall K.L.,

RA Achtman M., Akkin R., Baker S., Basham D., Bason N., Cherevach I.,

RA Achtman M., Atkin R., Baker S., Basham D., Bason N., Cherevach I.,

RA Achtman M., Atkin R., Baker S., Basham D., Bason N., Cherevach I.,

RA Chillingworth T., Collins M., Cronin A., Davis P., Doggett J.,

RA Feltwell T., Goble A., Hamlin N., Hauser H., Holroyd S., Jagels K.,

RA Leather S., Moule S., Norberczak H., O'Neil S., Ormond D., Price C.,

RA Leather S., Mutter S., Sanders M., Sanders D., Seeger K.,

RA Leather S., Simmonds M., Skelton J., Squares R., Squares S., Stevens K.,

RA Aharp S., Simmonds M., Skelton J., Squares R., Squares S., Stevens K.,

RA Mowin L., Whitehead S., Barrell B.G., Maskell D.J.;

"Comparative analysis of the genome sequences of Bordetella pertussis,

RT Mocdetella parapertussis and Bordetella bronchiseptica.";

RM Mat. Genet. 35:32-40(2003).

RM Phocomal Trotters. CAB43895.1;

RM Phocomal Trotters. CAB43895.1;
                                                                                            Matches
                                                                                                                Query Match
Best Local :
                                                                                                                                                                                   EMBL; J02
NON_TER
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Q9YPG8
Q9YPG8;
01-MAY-1999
                                                                                                                                                                                      phenotypes: Identification of 207-nucleotide product with protein kinase activity.";

J. Virol. 46:985-992(1983).
EMBL; J02349; AAA42577.1;
NON TER
40
3EQUENCE 40 77
                                                                                                                                                                                                                                                                                                                                                                                                                                                              01-MAY-1999 (TrEMBLrel. 10, Created)
01-MAY-1999 (TrEMBLrel. 10, Last sequence update)
01-MAY-1999 (TrEMBLrel. 10, Last annotation update)
(Prague 'wild type') src gene; amino acids 1-40 (Fragment).
Rous sarcoma virus.
Viruses; Retroid viruses; Retroviridae; Alpharetrovirus.
                                                                                                                                                                                                                                                                                                                            SEQUENCE FROM N.A.
MEDLINE=83216333; PubMed=6304353;
Kitamura N., Yoshida M.;
"Small deletion in src of Rous sa
                                                                                                                                                                                                                                                                                                                                                                                                                                            NCBI_TaxID=11886;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Ribosomal protein; Complete proteome.
SEQUENCE 37 AA; 4332 MW; 53B3EBBE5487DADD CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Bacteria; Proteobacteria; E
Alcaligenaceae; Bordetella.
NCBI_TaxID=520;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE FROM N.A.
STRAIN=Tohama I / ATCC BAA-589 / NCTC 13251;
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    7
                                            ω
                                                                                         Similarity 7; Conserv
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GPGDPGQRRRGLG
                                          GPTDPRHRSRNVG 21
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                                                                                                                                                                                   40 AA;
                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 30.9%; llarity 75.0%; Conservative
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                                                                                                             30.9%;
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                                                                                       Score 38; DB
Pred. No. 1.8e
1; Mismatches
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Pred. No. 1.7e
1; Mismatches
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                                                                                                             DB 15;
1.8e+02;
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1.7e+02;
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                                                                                                                                    Length 40;
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Q95WF4
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Best Local 9
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01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-JUN-2003 (TrEMBLrel. 24, Last sequence update)
1socytochrome c2 (cycl) (Fragment).
Rhodobacter sphaeroides (Rhodopseudomonas sphaeroides).
Bacteria; Proteobacteria; Alphaproteobacteria; Rhodobacterales;
Rhodobacteraceae; Rhodobacter.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           095WF4;
01-DEC-2001 (TrEMBLrel. 19, Created)
01-DEC-2001 (TrEMBLrel. 19, Last sequence update)
01-JUN-2003 (TrEMBLrel. 24, Last annotation updat
Nuclear receptor Amurlo (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PIR; C406
NON TER
SEQUENCE
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submitted (DEC-1992) to the
EMBL; L02104; AAA61342.1;
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Donohue T.J.;
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                                                                                                                                                                                                                                                                                                                                                                                                            MOI. Phylogenet. Bvol. 21:93-102(2001)
-!- SUBCELLULAR LOCATION: NUCLEAR (BY SIMILARITY)
-!- SIMILARITY: BELONGS TO THE NUCLEAR HORMONE RECEPTOR FAMIL)
EMBL; AF323690; AAL72203.1; --
GO; GO:0005634; C:nucleus; IEA.
GO; GO:0006872; F:receptor activity; IEA.
GO; GO:0004872; F:receptor activity; IEA.
GO; GO:0003700; F:transcription factor activity; IEA.
GO; GO:0006355; P:regulation of transcription, DNA-dependent;
GO; GO:0006350; P:transcription; IEA.
InterPro; IPR001628; Zaf_C4steroid.
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NON TER 1 1
NON TER 27 27
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MEDLINE=21490824; PubMed=11603940;
Grasso L.C., Hayward D.C., Trueman
Ball E.E.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Astrocoenina; Acroporidae;
NCBI_TaxID=45264;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Acropora millepora.
Eukaryota; Metazoa; Cnidaria; Anthozoa;
                                                                                                                                                                                                                                                   SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                     Pfam; PF00105; zf-C4;
DNA-binding; Metal-bii
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Acropora.";
Mol. Phylogenet.
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                                                                                                                        Similarity 6; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         QGFR PRHGATGDRHTLGLKRDRSV
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                                                                    PTDPRHRSR
      PVDKRHRNR
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                                                                                                                                                                                                                                                                                                                                 Metal-binding; Nucr
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRELIMINARY;
                                                                                                                              Conservative
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                                                                                                                                                                                                                                                   AA;
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3263 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          nuclear receptors:
                                                             18
      27
                                                                                                                                                                                                                                                   3302 MW;
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41.7%;
                                                                                                                                                     29.3%;
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to the EMBL/GenBank/DDBJ
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                                                                                                                                                                                                                                                                                                                                           Nuclear protein;
inc; Zinc-finger.
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                                                                                                                                                        Score 36;
Pred. No.
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Pred. No. 1
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                                                                                                                              Pred. No. 2.4
L; Mismatches
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PAC OCCUPANTO CONTROL OCCUPANT
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RESULT
Q7VKF4
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Matches 8
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Best Local S
Matches 7
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Q66546;
01-NOV-1996
01-NOV-1996
01-DEC-2001
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01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
EBV B95-8 Cl(e) DNA with antigen coding ORF (Fragment).
Human herpesvirus 4 (Epstein-Barr virus).
Viruses; dsDNA viruses, no RNA stage; Herpesviridae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE FROM N.A.

MEDLINE=88217505; PubMed=2835748;

Walls D., Perricaudet M., Gannon F.;

"The analysis of EBV proteins which a
Nucleic Acids Res. 16:289-2872(1988)

EMBL, X07530; CAA30405.1; -..

NON_TER
  Q7VKF4
Q7VKF4;
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01-MAR-2003
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NCBI_TaxID=10376;
                                                                                                                                                                                                                                                                                                                                                                  Derst C., Mescutat S., Walther C., Wicher D.;
"Cloning and Alternative Splicing of a large conductance activated potassium channel from Periplaneta americana.";
Submitted (NOV-2001) to the EMBL/GenBank/DDBJ databases.
EMBL; AF45216; F:ion channel activity; IEA.

GO; GO:0005216; F:ion channel activity; IEA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Periplaneta americana (American cockroach).
Eukaryota; Metazoa; Arthropoda; Hexapoda; Insecta;
Neoptera; Orthopteroidea; Dictyoptera; Blattaria; E
Blattidae; Periplaneta.
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01-JUN-2003
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Pred. No.
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NON TER
SEQUENCE
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STRAIN-35000HP / ATCC 700724;
MUNISON R.S. Jr., Ray W.C., Mahairas G., Sabo P., Mungur I Munson R.S. Jr., Ray W.C., Mahairas G., Hood L.;
"The complete genome sequence of Haemophilus ducreyi.";
Submitted (JUN-2003) to the EMBL/GenBank/DDBJ databases.
EMBL; A5017156; AAP96675.1;
Ribosomal protein; Complete proteome.
Ribosomal protein; Complete Schemer.
SEQUENCE 37 AA; 4279 MW; 53A83EB09A2F94ED CRC64;
                                                                                                                                                                                                                               SEQUENCE FROM N.A.

MEDLINE=95316863; PubMed=7796416;

Ji W., Qu G.Z., Ye P., Zhang X.Y., Halabi S., Ehrlich M.;

"Frequent detection of bcl-2/JH translocations in human blood and organ samples by a quantitative polymerase chain reaction assay.";

Cancer Res. 55:2876-2882 (1995).
                                                                                                                                                                                                                                                                                                                                                                                                                       01.WAY-1999 (TYEMBLIE) 10, Created)
01.WAY-1999 (TYEMBLIE) 10, Last seq
01.WAY-1999 (TYEMBLIE) 10, Last anno
10.WAY-1999 (TYEMBLIE) 10, Last anno
BC1-2-JH protein (Fragment).
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Bacteria; Proteobacteria; Gammaproteobacteria; Pasteurellales; Pasteurellaceae; Haemophilus.
NCBI_TaxID=730;
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505 ribosomal protein L36.
RPMJ2 OR HD1955.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Pred. No. 4.6e
2; Mismatches
                                                                                                                       Score 34.5; DB 4; Length 22; Pred. No. 3.2e+02; 1; Mismatches 3; Indels
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